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(54) Title: SUSCEPTIBILITY GENE FOR HUMAN STROKE; METHODS OF TREATMENT

(57) Abstract: A role of the human PDE4D gene in stroke is disclosed. Methods for diagnosis, prediction of clinical course and treatment for stroke using polymorphisms in the PDE4D gene are also disclosed.

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SUSCEPTIBILITY GENE FOR HUMAN STROKE;
METHODS OF TREATMENT

RELATED APPLICATIONS

5 This application is a continuation of and claims priority to U.S. Application No. 10/650,120, filed August 27, 2003, which is a continuation-in-part of U.S. Application No. 10/419,723 filed April 18, 2003, which is a continuation-in-part of U.S. Application No. 10/255,120, filed September 25, 2002, which is a continuation-in-part of U.S. Application No. 10/067,514, filed February 4, 2002, which is a
10 continuation-in-part of U.S. Application No. 09/811,352, filed March 19, 2001. The entire teachings of the above applications are incorporated herein by reference.

BACKGROUND OF THE INVENTION

15 Stroke is a common and serious disease. Each year in the United States more than 600,000 individuals suffer a stroke and more than 160,000 die from stroke-related causes (Sacco, R.L. *et al.*, *Stroke* 28, 1507-17 (1997)). In western countries stroke is the leading cause of severe disability and the third leading cause of death (Bonita, R., *Lancet* 339, 342-4 (1992)). The lifetime risk of those who reach the age of 40 exceeds 10%.

20 The clinical phenotype of stroke is complex but is broadly divided into ischemic (accounting for 80-90%) and hemorrhagic stroke (10-20%) (Caplan, L.R. *Caplan's Stroke: A Clinical Approach*, 1-556 (Butterworth-Heinemann, 2000)). Ischemic stroke is further subdivided into large vessel occlusive disease (referred to here as carotid stroke), usually due to atherosclerotic involvement of the common
25 and internal carotid arteries, small vessel occlusive disease, thought to be a non-atherosclerotic narrowing of small end-arteries within the brain, and cardiogenic stroke due to blood clots arising from the heart usually on the background of atrial fibrillation or ischemic (atherosclerotic) heart disease (Adams, H.P., Jr. *et al.*, *Stroke* 24, 35-41 (1993)). Therefore, it appears that stroke is not one disease but a
30 heterogeneous group of disorders reflecting differences in the pathogenic mechanisms (Alberts, M.J. *Genetics of Cerebrovascular Disease*, 386 (Futura

Publishing Company, Inc., New York, 1999); Hassan, A. & Markus, H.S. *Brain* 123, 1784-812 (2000)). However, all forms of stroke share risk factors such as hypertension, diabetes, hyperlipidemia, and smoking (Sacco, R.L. *et al.*, *Stroke* 28, 1507-17 (1997); Leys, D. *et al.*, *J. Neurol.* 249, 507-17 (2002)). Family history of stroke is also an independent risk factor suggesting the existence of genetic factors that may interact with environmental factors (Hassan, A. & Markus, H.S. *Brain* 123, 1784-812 (2000); Brass, L.M. & Alberts, M.J. *Baillieres Clin. Neurol.* 4, 221-45 (1995)).

The genetic determinants of the common forms of stroke are still largely unknown. There are examples of mutations in specific genes that cause rare Mendelian forms of stroke such as the *Notch3* gene in CADASIL (cerebral autosomal dominant arteriopathy with subcortical infarctions and leukoencephalopathy) (Tournier-Lasserre, E. *et al.*, *Nat. Genet.* 3, 256-9 (1993); Joutel, A. *et al.*, *Nature* 383, 707-10 (1996)), *Cystatin C* in the Icelandic type of hereditary cerebral hemorrhage with amyloidosis (Palsdottir, A. *et al.*, *Lancet* 2, 603-4 (1988)), *APP* in the Dutch type of hereditary cerebral hemorrhage (Levy, E. *et al.*, *Science* 248, 1124-6 (1990)) and the *KRIT1* gene in patients with hereditary cavernous angioma (Gunel, M. *et al.*, *Proc. Natl. Acad. Sci. USA* 92, 6620-4 (1995); Sahoo, T. *et al.*, *Hum. Mol. Genet.* 8, 2325-33 (1999)). None of these rare forms of stroke occur on the background of atherosclerosis, and therefore, the corresponding genes are not likely to play roles in the common forms of stroke which most often occur with atherosclerosis.

It is very important for the health care system to develop strategies to prevent stroke. Once a stroke happens, irreversible cell death occurs in a significant portion of the brain supplied by the blood vessel affected by the stroke. Unfortunately, the neurons that die cannot be revived or replaced from a stem cell population. Therefore, there is a need to prevent strokes from happening in the first place. Although we already know of certain clinical risk factors that increase stroke risk (listed above), there is an unmet medical need to define the genetic factors involved in stroke to more precisely define stroke risk. Further, if predisposing alleles are common in the general population and the specificity of predicting a disease based on their presence is low, additional loci such as protective loci are needed for

meaningful prediction of disposition of the disease state. There is also a great need for therapeutic agents for preventing the first stroke or further strokes in individuals who have suffered a previous stroke or transient ischemic attack.

5 SUMMARY OF THE INVENTION

A locus conferring susceptibility to ischemic stroke to chromosome 5q12 in the Icelandic population has been mapped and the identification of phosphodiesterase 4D (*PDE4D*) as the gene at 5q12 contributing to the risk of ischemic stroke has been reported. This locus was extensively fine mapped and tested for association to stroke.

10 Most striking is that haplotypes can be classified into three distinct groups: wild type, at-risk and protective. Additionally, a significant dysregulation of multiple *PDE4D* isoforms in stroke patients was observed. The strongest association was within the *PDE4D*, especially to the two major subtypes of ischemic stroke, carotid and cardiogenic stroke. We have found variation in *PDE4D* that more than doubles the risk for

15 cardiogenic and carotid stroke, two of the most common forms of ischemic stroke. We have shown that there are at least 9 isoforms of *PDE4D* at the mRNA level and the protein level. The basis for these isoforms is the use of alternative 5 prime exons that are alternatively spliced into a common set of exons defining the catalytic domain as well as, in the case of the long forms, a set of exons defining a common core in the regulatory

20 domain. The *PDE4D* gene is involved in the pathogenesis of stroke. The *PDE4D* gene may be involved through atherosclerosis, the major pathological process underlying ischemic stroke. Our results indicate that atherosclerosis is a cAMP disease resulting from dysregulation of its levels within the vasculature.

In one aspect, the invention relates to methods of diagnosing a predisposition

25 to stroke. The methods of diagnosing a predisposition to stroke in an individual include detecting the presence of a polymorphism in *PDE4D*, as well as detecting alterations in expression of a *PDE4D* polypeptide or isoform, such as the presence of, or relative expression of different splicing variants of *PDE4D* polypeptides. For example, it may be that the ratio of certain splice variants could be used as a

30 diagnostic marker for stroke predisposition. Also an abnormal splice form can be detected (that is one that is not normally expressed but is created from a DNA sequence mutation that leads to an abnormal splice form to be created from the primary transcript) may be created from mutations in the *PDE4D* gene. For example,

new splice sites might be created from a single base substitution within an intron that is inappropriately used as a splice acceptor or donor site, resulting in an abnormal message which is likely to have a premature stop codon leading to a truncated form of PDE4D protein. The alterations in expression can be quantitative, qualitative, or both quantitative and qualitative. The methods of the invention allow the accurate diagnosis of stroke at or before disease onset, thus reducing or minimizing the debilitating effects of stroke. The methods of the invention also diagnose those individuals who are protected against developing stroke even in the face of other risk factors including but not restricted to hypertension, diabetes, hyperlipidemia, smoking history, previous stroke, TIA, MI or PAOD, or carriers of stroke associated gene variants. In one embodiment, predisposition to stroke or susceptibility to stroke can be assessed by determining PDE4D isoform levels in the individual compared to control levels, wherein a difference in isoform expression is indicative of predisposition or susceptibility to stroke. Preferably, the level of expression of PDE4D7 and/or PDE4D9 is assessed.

The invention additionally relates to an assay for identifying agents that alter (*e.g.*, enhance or inhibit) the activity or expression or transcription of one or more PDE4D polypeptides or isoforms. Such an assay may also identify agents that alter the relative expression of one or more PDE4D isoforms with respect to other isoforms at either the mRNA level or polypeptide level. For example, a cell, cellular fraction, or solution containing a PDE4D polypeptide or a fragment or derivative thereof, can be contacted with an agent to be tested, and the level of PDE4D polypeptide expression or activity can be assessed. Alternatively, a cell, or cell with artificial DNA construct with part or all of the PDE4D gene with or without a reporter gene can be used to identify agents that may directly affect transcription at one or more of the many alternative PDE4D promoters upstream of the alternative 5 prime exons or splicing efficiency of the primary transcript to one or more mRNA isoforms. The activity or expression of more than one PDE4D polypeptides can be assessed concurrently (or the corresponding reporter gene activity) (*e.g.*, the cell, cellular fraction, or solution can contain more than one type of PDE4D polypeptide, such as different splicing variants, and the levels of the different polypeptides or splicing mRNA variants can be assessed).

Agents that enhance or inhibit PDE4D mRNA or polypeptide expression or activity are also included in the current invention, as are methods of altering (enhancing or inhibiting) PDE4D mRNA or polypeptide expression or activity by contacting a cell containing PDE4D gene, mRNA, and/or polypeptide, or by
5 contacting the PDE4D gene, mRNA, and/or polypeptide, with an agent that enhances or inhibits expression or activity of PDE4D mRNA or polypeptide. In another embodiment, isoform mRNA and/or protein levels can be altered, compared to control levels, using the agents of the invention.

Additionally, the invention pertains to pharmaceutical compositions
10 comprising the nucleic acids of the invention, the polypeptides of the invention, and/or the agents that alter activity of PDE4D polypeptide. The invention further pertains to methods of treating stroke, by administering PDE4D therapeutic agents, such as nucleic acids of the invention, polypeptides of the invention, the agents that alter activity of PDE4D polypeptide, or compositions comprising the nucleic acids,
15 polypeptides, and/or the agents that alter activity of PDE4D polypeptide.

The invention further relates to methods for preventing the occurrence of stroke in an individual in need thereof by regulating a PDE4D mRNA and/or polypeptide isoform level compared to control levels, whereby the regulated isoform level mimics the level of a healthy individual. Isoform expression at the mRNA
20 and/or polypeptide level can be regulated using the agents and pharmaceutical compositions of the invention, by genetic alteration, by altering the ratio of isoforms and/or their absolute expression. In one embodiment, isoforms PDE4D7 and/or PDE4D9 can be regulated.

The invention further provides a method of diagnosing susceptibility to stroke
25 in an individual. This method comprises screening for one of the at-risk haplotypes in the phosphodiesterase 4D gene that is more frequently present in an individual susceptible to stroke, compared to the frequency of its presence in the general population, wherein the presence of an at-risk haplotype is indicative of a susceptibility to stroke. An "at-risk haplotype" is intended to embrace one or a
30 combination of haplotypes described herein over the PDE4D gene that show high correlation to stroke. In one embodiment, the at-risk haplotype is characterized by the presence of at least one single nucleotide polymorphism at nucleic acid positions

at risk haplotype 1 is G at nucleic acid position 142780 respectively, relative to SEQ ID NO: 1 and allele 0 of microsatellite marker AC0088181-1. In another embodiment, the at-risk haplotype 2 is characterized by the presence of at least one single nucleotide polymorphism and microsatellite marker at nucleic acid positions
5 142780, 135112, 132562, 131865, 129361, 129360, 125304, 123426, 123312, 120628, 118914, 111781, 111252, 109301, 107849, 105225, 104552, 102977, 100795, 99035, 88614, 88456, 83119, 82244, 80127, 78552, relative to SEQ ID NO: 1 and allele 0 microsatellite marker AC0088181-1.

In yet another embodiment, the at-risk haplotype 3 is characterized by the
10 presence of at least one polymorphism at nucleic acid positions 138806, 131865, 129361, 120628, 91470 relative to SEQ ID NO: 1.

Also described are methods for diagnosing susceptibility to stroke in an individual comprising screening for an at-risk haplotype in the phosphodiesterase 4D gene that is more frequently present in an individual susceptible to stroke (affected),
15 compared to the frequency of its presence in a healthy individual (control) wherein the screening for the presence of an at-risk haplotype within or near PDE4D that significantly correlates with at least one of the haplotypes described herein or stroke susceptibility. As an example of a simple test for correlation would be a Fisher-exact test on a two by two table. Given a cohort of chromosomes the two by two table is
20 constructed out of the number of chromosomes that include both of the haplotypes, one of the haplotype but not the other and neither of the haplotypes.

A protective haplotype is intended to embrace one or a combination of haplotypes described herein over the PDE4D gene that show a protective characteristic or property of a reduced risk of stroke. The particular combination of
25 genetic markers (haplotypes) are present at a higher than expected frequency in controls than patients. Individuals with a protective allele or haplotype are about 30% less likely to have a stroke compared to the general population. In one embodiment, a protective haplotype is characterized by the presence of at least one single nucleotide polymorphism, such as the allele A at nucleotide position 142780
30 relative to SEQ ID NO: 1. The presence of the polymorphisms that comprise the at-risk haplotype or protective haplotype can be determined by electrophoretic analysis, restriction length polymorphism analysis, fluorescence energy transfer detection,

kinetic PCR, allele specific PCR, sequence analysis, hybridization analysis or other known techniques.

Kits for diagnosing susceptibility to stroke in an individual are also disclosed and comprise primers for nucleic acid amplification of a region of PDE4D
5 comprising the at-risk haplotype and/or protective haplotype.

The first major application of the current invention involves prediction of those at higher risk of developing a stroke. Diagnostic tests that define genetic factors contributing to stroke might be used together with or independent of the known clinical risk factors to define an individual's risk relative to the general
10 population. Better means for identifying those individuals at risk for stroke should lead to better prophylactic and treatment regimens, including more aggressive management of the current clinical risk factors such as hypertension, diabetes, hypercholesterolemia, hypertriglyceridemia, obesity, and inflammatory components as reflected by increased C-reactive protein levels or other inflammatory markers.
15 Information on genetic risk may be used by physicians to help convince particular patients to adjust life style and quit smoking. This invention provides the means to define a genetic component that doubles an individual's risk for stroke. Also described are means to define the genetic components that protect an individual from stroke.

20 The second major application of the current invention is the specific identification of a rate-limiting pathway involved in stroke. While many have attempted to find genes that are over-expressed or under-expressed in atherosclerosis plaques in the carotid arteries, the vast majority of the changes seen in diseased blood vessels compared to normal blood vessels are simply a reaction to the underlying
25 process of atherosclerosis and stroke predisposition and are not the underlying cause. A disease gene with genetic variation that is significantly more common in stroke patients as compared to controls represents a specifically validated causative step in the pathogenesis of stroke. That is, the uncertainty about whether a gene is causative or simply reactive to the disease process is eliminated. The protein encoded by the
30 disease gene defines a rate-limiting molecular pathway involved in the biological process of stroke predisposition. The proteins encoded by such stroke genes or its interacting proteins in its molecular pathway may represent drug targets that may be

selectively modulated by small molecule, protein, antibody, or nucleic acid therapies. Such specific information is greatly needed since stroke prevention and treatment is a major unmet medical need that affects over a half-million Americans each year.

Also useful is determining the gene that is protective against stroke. The proteins
5 encoded by the protective gene and the biological pathway that it is a member may represent another target selectively modulated by small molecule, protein antibody or nucleic acid therapies.

A third application of the current invention is its use to predict an individual's response to a particular drug, even drugs that do not act on PDE4D or its pathway. It
10 is a well-known phenomenon that in general, patients do not respond equally to the same drug. Much of the differences in drug response to a given drug is thought to be based on genetic and protein differences among individuals in certain genes and their corresponding pathways. Our invention defines the PDE4D pathway and its effect on cAMP levels in cells where it is expressed as one key molecular pathway
15 involved in stroke risk. Some current or future therapeutic agents may be able to affect this pathway directly or indirectly and therefore, be effective in those patients whose stroke risk is in part determined by PDE4D pathway genetic variation. On the other hand, those same drugs may be less effective or ineffective in those patients who do not have at risk variation in the PDE4D gene or pathway. Therefore, PDE4D
20 variation or haplotypes may be used as a pharmacogenomic diagnostic to predict drug response and guide choice of therapeutic agent in a given individual.

The invention helps meet the unmet medical needs in at least two major ways: 1) it provides a means to define patients at higher risk for stroke than the general population who can be more aggressively managed by their physicians in an
25 effort to prevent stroke; and 2) it defines a drug target that can be used to screen and develop therapeutic agents that can be used to prevent stroke before it happens or prevent a second stroke in those who have already suffered a stroke or transient ischemic attack.

BRIEF DESCRIPTION OF THE DRAWINGS

The foregoing and other objects, features and advantages of the invention will be apparent from the following more particular description of certain embodiments of the invention, as illustrated in the accompanying drawings.

5 FIGS. 1.1 and 1.2 show two family pedigrees each affected by several of the stroke subtypes, including hemorrhagic stroke.

FIGS. 2.1, 2.2 and 2.3 show the genetic, combined and physical maps for locating the PDE4D gene using 30 polymorphic markers. For the combined map, all markers have been assigned in the genetic and physical map unless otherwise
10 indicated (* indicates marker only assigned in the physical map; ** indicates markers only assigned in genetic map).

FIG. 3 shows the schematic representations of PDE4D splice variants. Splice variants PDE4D9 are novel, as well as exons D7A-1, D7A-2, D7A-3, D8 and D9. Splice variants 4DN1, 4DN2 and 4DN3 (Miro, *et al.*, *Biochem. Biophys. Res.*
15 *Comm.*, 274: 415-421 (2002), and 4D1, 4D2, 4D3, 4D4 and 4D5 are known (Bolger *et al.*, *Biochem. J.* pt. 2: 539-548 (1997)).

FIG. 4 is a graphic representation showing PDE4D isoform expression in EBV transformed cells (expression of PDE4D3 and PDE4D9 below detection limits).

FIG. 5 is a graphic representation showing expression of PDE4D isoforms in
20 EBV transformed cells from patients with or without the stroke-associated haplotype.

FIG. 6 is a graphic representation showing expression of PDE4D isoforms in EBV cells from controls with or without the stroke-associated haplotype.

FIGS. 7.1 to 7.10 show the amino acid sequences for the isoforms of the PDE4D gene. SEQ ID NO: 2 is D4; SEQ ID NO: 3 is N2; SEQ ID NO: 4 is D5;
25 SEQ ID NO: 5 is N3; SEQ ID NO: 6 is D3; SEQ ID NO: 7 is N1; SEQ ID NO: 8 is D8; SEQ ID NO: 9 is D1; and SEQ ID NO: 10 is D2.

FIGS. 8.1 and 8.2 list all publicly available PDE4D mRNAs and novel cDNA segments identified by deCODE genetics.

FIGS. 9.1 to 9.351 show the genomic sequence of the human PDE4D gene.

30 FIGS. 10.1 to 10.3 show a graphic representation showing the single marker allelic association within the PDE4D gene. FIG. 10.1 is a schematic showing the gene structures. FIG. 10.2 shows graphic representation of the microsatellite and

SNP distribution within the *PDE4D* gene. FIG. 10.3 shows graphic representation of the single marker allelic association across the *PDE4D* gene for both microsatellites (filled circles) and SNPs (open circles); negative log p-value versus the physical location in kilobases.

5 FIGS. 11.1 to 11.3 graphically depict the haplotype association for carotid and cardiogenic stroke combined. Estimated haplotype frequencies for patients and controls respectively, are indicated within parentheses. FIG. 11.1 is a comparison of groups of haplotypes constructed from SNP45 and AC008818-1, two markers separated by 6kb. Note that X is a *composite* allele that denotes jointly all alleles of
10 AC008818-1 except allele 0. Apart from haplotype A0 that is not found in our samples, other haplotypes can be grouped into three groups with distinct risks. Each arrow corresponds to a comparison between two groups and RR is the estimated risk of the group the arrow is pointing at relative to the other group. The difference between 1 and the information (Info) is a measure of the fraction of information that
15 is lost due to uncertainty in phase and missing genotypes. FIG. 11.2 shows intermediate results when the investigation is extended from SNP45 and AC008818-1, which are both in LD block B, to include 25 SNPs in LD block C. H_C is the at-risk haplotype, identified in FIG. 13 and L_C is a composite haplotype that denotes jointly all haplotypes of the 25 SNPs except H_C . Together with AC008818-1 and
20 SNP45, the haplotypes here span 64kb. Haplotype G0 in A is split into extended haplotypes $G0H_C$ and $G0L_C$. $G0H_C$ has significantly higher risk than $G0L_C$, and the risk of $G0L_C$ is not distinguishable from the wild type GX. FIG. 11.3 shows a refinement of the groupings in A — $G0L_C$ is moved from the at-risk group to the wild type group. Also noted is that the extended haplotype AXH_C does not exist
25 indicating that blocks B and C are in LD.

FIG. 12 is a schematic representation of the physical map of *STRK1* interval showing all genes and mRNAs in region. Markers identified with an asterisk (*) indicate those with significant single marker association.

30 FIGS 13.1 to 13.3 show a graphical depiction of the linkage disequilibrium (LD) and haplotypes in the 5' end of *PDE4D* gene. FIG. 13.1 shows pairwise linkage disequilibrium between SNPs in a 600 kb region in the 5' end of *PDE4D*. The markers are plotted equidistant. Two measures of LD are shown: D' in the upper left

triangle and p-values in the lower right triangle. This region can be divided into three blocks of strong LD, each with limited haplotype diversity, block A, block B and block C. The lines indicate the position of the three exons D7-1, D7-2 and D7-3 and the microsatellite marker AC008818-1. FIG. 13.2 show all common haplotypes
5 identified within each of the three blocks. Association results for all the haplotypes are presented in Table 2C. FIG. 13.3 depicts the percentage of chromosomes within each block that match one of the common haplotypes.

DETAILED DESCRIPTION OF THE INVENTION

10 The first major stroke locus, *STRK1*, was mapped to 5q12 using a genome-wide search for susceptibility genes in the common forms of stroke. A broad but rigorous definition of the phenotype was used including patients with ischemic stroke, transient ischemic attack (TIA), and hemorrhagic stroke. The lod score after adding a higher density of markers (one marker every 1 cM) was 4.40 ($P=3.9 \times 10^{-6}$)
15 at marker D5S2080. The lod score increased to 4.9 after the hemorrhagic stroke patients were removed, suggesting that the gene at the locus is primarily important for ischemic stroke. The most promising region harboring a stroke susceptibility gene was narrowed down to a segment less than 6 cM (approximately 3.8 Mb), from D5S1474 to D5S398, as defined by a decrease of one in LOD score (will be referred
20 to as the "one-LOD interval" hereafter).

We describe here the positional cloning of a stroke susceptibility gene located in the *STRK1* locus. This region was extensively fine-mapped and tested for association to stroke. The strongest association found in the one-LOD interval was within the phosphodiesterase 4D gene (*PDE4D*), a member of the large superfamily
25 of cyclic nucleotide phosphodiesterases. The strongest signal observed at *PDE4D* was to the two major subtypes of ischemic stroke, carotid and cardiogenic stroke. Relative expression of *PDE4D* isoforms correlated with stroke and with the genetic variation within *PDE4D* which is associated to stroke. Our results suggest that this gene is involved in pathogenesis of stroke through atherosclerosis, the major
30 pathological process underlying stroke.

Our results also indicate that genetic variation in the *PDE4D* gene is associated with ischemic stroke. The direct involvement of *PDE4D* is strongly supported by both

linkage and haplotype association. Multiple markers and haplotypes within the *PDE4D* gene show strong association to stroke. The haplotypes can be classified into three distinct groups, wild type, at-risk and protective. We first identified the association using microsatellite markers, and supplementing the microsatellite data with a denser set of SNPs further supported this. The strongest association was to the two ischemic subtypes, carotid and cardiogenic stroke. This gene shows no association to small vessel occlusive disease, the form of stroke thought to be independent of atherosclerosis. Haplotype analyses show that the most significant haplotype extends over an area of 260 kb covering the first exon of the *PDE4D* gene. The haplotype is significantly associated to carotid and cardiogenic stroke with a relative risk of 2.3 and approximately 47 % of carotid/cardiogenic stroke patients carry at least one copy of this haplotype. This same haplotype has a relative risk of 1.8 for stroke in general. This haplotype extends over the 5' exon unique to the *PDE4D7* isoform and the presumed promoter region of this isoform suggesting that the functional variation may be involved in transcriptional regulation. This hypothesis is also supported by our *PDE4D* expression analysis that shows that there is significant correlation between the disease associated haplotype and the level of *PDE4D7* message.

The strongest association found for this *PDE4D* haplotype was to the two major subtypes of ischemic stroke, carotid and cardiogenic stroke suggesting a role for this gene in the vascular biology of atherosclerosis. While there are multiple etiologies for ischemic stroke, atherosclerosis remains the most important one. Atherosclerosis is a chronic progressive disease characterized by accumulation of lipids, fibrous, and cellular elements within the large arteries. These lesions can grow sufficiently large to impede blood flow and, more importantly, their surfaces can rupture leading to local thrombus formation occluding the blood vessel and causing a stroke or myocardial infarction. The major pathological process for the two ischemic subtypes, carotid and cardiogenic stroke is atherosclerosis. First, it is the major cause of stenotic and occlusive lesions of the internal and common carotids that lead to carotid strokes. Second, cardiac thrombi which shed emboli to the brain most commonly occur on the background of coronary artery disease, such as following acute myocardial infarction or ischemic cardiomyopathy, and/or due to atrial fibrillation on the basis of poor compliance of ischemic ventricles (diastolic dysfunction/stiffening). Although atrial fibrillation may occur on the background of

other diseases such as valvular disease, hyperthyroidism, and hypertension, in the age group that tends to suffer from stroke, ischemic heart disease remains one of the most important causes. Ischemic stroke resulting from occlusion of small penetrating arteries within the brain (small vessel occlusive disease or lacunar stroke) is generally thought to result from local endothelial proliferation since atherosclerosis only occurs in larger arteries. PDE4D does not show association to small vessel stroke, consistent with its role in atherosclerosis. In summary, atherosclerosis accounts for the majority of all strokes, particularly carotid and cardiogenic stroke, two subphenotypes that show the strongest association to the *PDE4D* gene.

REPRESENTATIVE TARGET POPULATION

An individual at risk for stroke is an individual who has at least one risk factor, such as previous stroke or TIA, an at-risk haplotype in one or more stroke risk genes, an at-risk haplotype for the *PDE4D* gene; a polymorphism in a *PDE4D* gene; dysregulation of *PDE4D* isoform expression; diabetes; hypertension; hypercholesterolemia; elevated lip(a); obesity; a past or current smoker; an elevated inflammatory marker (e.g., a marker such as C-reactive protein (CRP), serum amyloid A, fibrinogen, tissue necrosis factor-alpha, a soluble vascular cell adhesion molecule (sVCAM), a soluble intervascular adhesion molecule (sICAM), E-selectin, matrix metalloproteinase type-1, matrix metalloproteinase type-2, matrix metalloproteinase type-3, and matrix metalloproteinase type-9); increased LDL cholesterol and/or decreased HDL cholesterol; and/or at least one previous myocardial infarction, concurrent MI, acute coronary syndrome, stable angina, atherosclerosis, carotid stenosis, peripheral vascular occlusive disease, or requires treatment for restoration of coronary artery blood flow (e.g., angioplasty, stent, coronary artery bypass graft).

An individual who has a protective haplotype is one who is less likely to have a stroke. In another embodiment of the invention, an individual who is at risk for stroke is an individual who has a polymorphism in a *PDE4D* gene, in which the presence of the polymorphism is indicative of a susceptibility to stroke. An individual who has a protective haplotype and less likely to have a stroke is an individual who has a polymorphism in a *PDE4D* gene such as the A allele at

nucleotide position 142780 relative to SEQ ID NO: 1, in which the presence of the polymorphism is indicative of a protection from stroke. The term "gene," as used herein, refers to not only the sequence of nucleic acids encoding a polypeptide, but also the promoter regions, transcription enhancement elements, splice donor/acceptor sites, splice enhancer and silencer sequences and other regulators of splicing, and other non-transcribed nucleic acid elements. Representative polymorphisms include those presented in Table 11, below.

In one embodiment of the invention, an individual who is at risk for stroke is an individual who has an at-risk haplotype in PDE4D, as described herein, particularly but not limited to ischemic stroke. Increased risk for the two major subtypes of ischemic stroke, carotid and cardiogenic stroke, can be assessed by screening for at-risk haplotype that comprises SNP5PDM361194, SNP5PDM368135, SNP5PDM370640, SNP5PDM379372 and SNP5PDM408531 at the 5' UTR of PDE4D7. Results reported herein indicate that PDE4D is involved in pathogenesis of stroke through atherosclerosis. The major pathological process for carotid stroke and cardiogenic stroke is atherosclerosis. Thus, an individual who is at-risk for atherosclerosis, peripheral arterial occlusive disease, or myocardial infarction can also benefit from the teachings of the invention.

ASSESSMENT FOR AT-RISK AND PROTECTIVE HAPLOTYPES

A "haplotype," as described herein, refers to a combination of genetic markers ("alleles"), such as those set forth in Tables 1, 2C, 4A and 4B. In a certain embodiment, the haplotype can comprise one or more alleles, two or more alleles, three or more alleles, four or more alleles, or five or more alleles. The genetic markers are particular "alleles" at "polymorphic sites" associated with PDE4D. A nucleotide position at which more than one sequence is possible in a population (either a natural population or a synthetic population, *e.g.*, a library of synthetic molecules), is referred to herein as a "polymorphic site". Where a polymorphic site is a single nucleotide in length, the site is referred to as a single nucleotide polymorphism ("SNP"). For example, if at a particular chromosomal location, one member of a population has an adenine and another member of the population has a thymine at the same position, then this position is a polymorphic site, and, more

specifically, the polymorphic site is a SNP. Polymorphic sites can allow for differences in sequences based on substitutions, insertions or deletions. Each version of the sequence with respect to the polymorphic site is referred to herein as an “allele” of the polymorphic site. Thus, in the previous example, the SNP allows for
5 both an adenine allele and a thymine allele.

Typically, a reference sequence is referred to for a particular sequence. Alleles that differ from the reference are referred to as “variant” alleles. For example, the reference PDE4D sequence is described herein by SEQ ID NO: 1. The term, “variant PDE4D”, as used herein, refers to a sequence that differs from SEQ ID
10 NO: 1, but is otherwise substantially similar. The genetic markers that make up the haplotypes described herein are PDE4D variants.

Additional variants can include changes that affect a polypeptide, *e.g.*, the PDE4D polypeptide. These sequence differences, when compared to a reference nucleotide sequence, can include the insertion or deletion of a single nucleotide, or of
15 more than one nucleotide, resulting in a frame shift; the change of at least one nucleotide, resulting in a change in the encoded amino acid; the change of at least one nucleotide, resulting in the generation of a premature stop codon; the deletion of several nucleotides, resulting in a deletion of one or more amino acids encoded by the nucleotides; the insertion of one or several nucleotides, such as by unequal
20 recombination or gene conversion, resulting in an interruption of the coding sequence of a reading frame; duplication of all or a part of a sequence; transposition; or a rearrangement of a nucleotide sequence, as described in detail above. Such sequence changes alter the polypeptide encoded by a PDE4D nucleic acid. For example, if the change in the nucleic acid sequence causes a frame shift, the frame
25 shift can result in a change in the encoded amino acids, and/or can result in the generation of a premature stop codon, causing generation of a truncated polypeptide. Alternatively, a polymorphism associated with stroke or a susceptibility to stroke can be a synonymous change in one or more nucleotides (*i.e.*, a change that does not result in a change in the amino acid sequence). Such a polymorphism can, for
30 example, alter splice sites, affect the stability or transport of mRNA, or otherwise affect the transcription or translation of the polypeptide. The polypeptide encoded by the reference nucleotide sequence is the “reference” polypeptide with a particular

reference amino acid sequence, and polypeptides encoded by variant alleles are referred to as "variant" polypeptides with variant amino acid sequences.

Haplotypes are a combination of genetic markers, *e.g.*, particular alleles at polymorphic sites. The haplotypes described herein, *e.g.*, having markers such as
5 those shown in Table 3, Table 4A and 4B, are found more frequently in individuals with stroke than in individuals without stroke. Therefore, these haplotypes have predictive value for detecting stroke or a susceptibility to stroke in an individual. The haplotypes described herein are a combination of various genetic markers, *e.g.*, SNPs and microsatellites. Therefore, detecting haplotypes can be accomplished by
10 methods known in the art for detecting sequences at polymorphic sites, such as the methods described above.

In certain methods described herein, an individual who is at risk for stroke is an individual in whom an at-risk haplotype is identified. In one embodiment, the at-risk haplotype is one that confers a significant risk of stroke. In one embodiment,
15 significance associated with a haplotype is measured by an odds ratio. In a further embodiment, the significance is measured by a percentage. In one embodiment, a significant risk is measured as an odds ratio of at least about 1.2, including but not limited to: 1.2, 1.3, 1.4, 1.5, 1.6, 1.7, 1.8 and 1.9. In a further embodiment, an odds ratio of at least 1.2 is significant. In a further embodiment, an odds ratio of at least
20 about 1.5 is significant. In a further embodiment, a significant increase in risk is at least about 1.7 is significant. In a further embodiment, a significant increase in risk is at least about 20%, including but not limited to about 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95% and 98%. In a further embodiment, a significant increase in risk is at least about 50%. It is understood
25 however, that identifying whether a risk is medically significant may also depend on a variety of factors, including the specific disease, the haplotype, and often, environmental factors.

An at-risk haplotype in, or comprising portions of, the PDE4D gene, is one where the haplotype is more frequently present in an individual at risk for stroke
30 (affected), compared to the frequency of its presence in a healthy individual (control), and wherein the presence of the haplotype is indicative of stroke or susceptibility to stroke. A protective haplotype in or comprising portions of the

PDE4D gene is one where the haplotype is more frequently present in an individual where the haplotype is protective against being affected by stroke compared to the frequency of its presence in an individual with stroke. The presence of the haplotype is indicative of a protection from stroke or protection from susceptibility to stroke as described above.

Standard techniques for genotyping for the presence of SNPs and/or microsatellite markers can be used, such as fluorescent-based techniques (Chen, *et al.*, *Genome Res.* 9, 492 (1999)), PCR, LCR, Nested PCR and other techniques for nucleic acid amplification. In one embodiment, the method comprises assessing in an individual the presence or frequency of SNPs and/or microsatellites in, comprising portions of, the PDE4D gene, wherein an excess or higher frequency of the SNPs and/or microsatellites compared to a healthy control individual is indicative that the individual has stroke, or is susceptible to stroke. See, for example, Table 1, Table 2C, Table 2D, Table 3, Table 4A and 4B (below) for SNPs and markers that can form haplotypes that can be used as screening tools. These markers and SNPs can be identified in at-risk haplotypes. For example, an at-risk haplotype can include microsatellite markers and/or SNPs such as those set forth in Table 2C, Table 4B and 4B. The presence of the haplotype is indicative of stroke, or a susceptibility to stroke, and therefore is indicative of an individual who falls within a target population for the treatment methods described herein.

Haplotype analysis first involves defining a candidate susceptibility locus using LOD scores. The defined regions are then ultra-fine mapped with microsatellite markers with an average spacing between markers of less than 100kb. All usable microsatellite markers that found in public databases and mapped within that region can be used. In addition, microsatellite markers identified within the deCODE genetics sequence assembly of the human genome can be used. The frequencies of haplotypes in the patient and the control groups using an expectation-maximization algorithm can be estimated (Dempster A. *et al.*, 1977. *J. R. Stat. Soc. B*, 39:1-389). An implementation of this algorithm that can handle missing genotypes and uncertainty with the phase can be used. Under the null hypothesis, the patients and the controls are assumed to have identical frequencies. Using a likelihood approach, an alternative hypothesis where a candidate at-risk-haplotype,

which can include the markers described herein, is allowed to have a higher frequency in patients than controls, while the ratios of the frequencies of other haplotypes are assumed to be the same in both groups is tested. Likelihoods are maximized separately under both hypotheses and a corresponding 1-df likelihood ratio statistics is used to evaluate the statistic significance.

To look for at-risk-haplotypes in the 1-lod drop or protective haplotypes, for example, association of all possible combinations of genotyped markers is studied, provided those markers span a practical region. The combined patient and control groups can be randomly divided into two sets, equal in size to the original group of patients and controls. The haplotype analysis is then repeated and the most significant p-value registered is determined. This randomization scheme can be repeated, for example, over 100 times to construct an empirical distribution of p-values.

In one embodiment, the at-risk haplotype is characterized by the presence of the polymorphism(s) represented by one or a combination of single nucleotide polymorphisms at nucleic acid positions 1425923, 1415979, 1414804, 1371388, 1307403 and 1257206, relative to SEQ ID NO: 1. In another embodiment, a diagnostic method for susceptibility to stroke can comprise determining the presence of at-risk haplotype represented by one or a combination of single nucleotide polymorphisms and microsatellite markers at nucleic acid positions 263539, 252772, 189780, 175259, 171240, 136550 and 120628, relative to SEQ ID NO: 1. In another embodiment, the at-risk haplotype is characterized by the following SNPs: SNP5PDM361194, SNP5PDM368135, SNP5PDM370640, SNP5PDM379372, and SNP5PDM408531. In one embodiment, the protective haplotype comprises the A allele of SNP45 at position 142780 relative to SEQ ID NO: 1. This haplotype is particularly useful for assessing susceptibility to the two major subtypes of ischemic stroke, carotid and cardiogenic stroke. In another embodiment, an at-risk haplotype, particularly for carotid and cardiogenic stroke, is characterized by use of microsatellite marker AC008818-1 to define the presence of an at-risk allele.

NUCLEIC ACID THERAPEUTIC AGENTS

In another embodiment, a nucleic acid of the invention; a nucleic acid complementary to a nucleic acid of the invention; or a portion of such a nucleic acid (e.g., an oligonucleotide as described below); or a nucleic acid encoding a PDE4D polypeptide, can be used in "antisense" therapy, in which a nucleic acid (e.g., an oligonucleotide) which specifically hybridizes to the mRNA and/or genomic DNA of a nucleic acid is administered or generated *in situ*. The antisense nucleic acid that specifically hybridizes to the mRNA and/or DNA inhibits expression of the polypeptide encoded by that mRNA and/or DNA, e.g., by inhibiting translation and/or transcription. Binding of the antisense nucleic acid can be by conventional base pair complementarity, or, for example, in the case of binding to DNA duplexes, through specific interaction in the major groove of the double helix.

An antisense construct can be delivered, for example, as an expression plasmid as described above. When the plasmid is transcribed in the cell, it produces RNA that is complementary to a portion of the mRNA and/or DNA that encodes a PDE4D polypeptide. Alternatively, the antisense construct can be an oligonucleotide probe that is generated *ex vivo* and introduced into cells; it then inhibits expression by hybridizing with the mRNA and/or genomic DNA of the polypeptide. In one embodiment, the oligonucleotide probes are modified oligonucleotides that are resistant to endogenous nucleases, e.g., exonucleases and/or endonucleases, thereby rendering them stable *in vivo*. Exemplary nucleic acid molecules for use as antisense oligonucleotides are phosphoramidate, phosphothioate and methylphosphonate analogs of DNA (see also U.S. Patent Nos. 5,176,996, 5,264,564 and 5,256,775). Additionally, general approaches to constructing oligomers useful in antisense therapy are also described, for example, by Van der Krol *et al.* (*Biotechniques* 6:958-976 (1988)); and Stein *et al.* (*Cancer Res.* 48:2659-2668 (1988)). With respect to antisense DNA, oligodeoxyribonucleotides derived from the translation initiation site are preferred.

To perform antisense therapy, oligonucleotides (mRNA, cDNA or DNA) are designed that are complementary to mRNA encoding the polypeptide. The antisense oligonucleotides bind to mRNA transcripts and prevent translation. Absolute complementarity, although preferred, is not required. A sequence "complementary"

to a portion of an RNA, as referred to herein, indicates that a sequence has sufficient complementarity to be able to hybridize with the RNA, forming a stable duplex; in the case of double-stranded antisense nucleic acids, a single strand of the duplex DNA may thus be tested, or triplex formation may be assayed. The ability to
5 hybridize will depend on both the degree of complementarity and the length of the antisense nucleic acid, as described in detail above. Generally, the longer the hybridizing nucleic acid, the more base mismatches with an RNA it may contain and still form a stable duplex (or triplex, as the case may be). One skilled in the art can ascertain a tolerable degree of mismatch by use of standard procedures.

10 The oligonucleotides used in antisense therapy can be DNA, RNA, or chimeric mixtures or derivatives or modified versions thereof, single-stranded or double-stranded. The oligonucleotides can be modified at the base moiety, sugar moiety, or phosphate backbone, for example, to improve stability of the molecule, hybridization, etc. The oligonucleotides can include other appended groups such as
15 peptides (*e.g.* for targeting host cell receptors *in vivo*), or agents facilitating transport across the cell membrane (see, *e.g.*, Letsinger *et al.*, *Proc. Natl. Acad. Sci. USA* 86:6553-6556 (1989); Lemaitre *et al.*, *Proc. Natl. Acad. Sci. USA* 84:648-652 (1987); PCT International Publication No. WO 88/09810) or the blood-brain barrier (see, *e.g.*, PCT International Publication No. WO 89/10134), or hybridization-triggered
20 cleavage agents (see, *e.g.*, Krol *et al.*, *BioTechniques* 6:958-976 (1988)) or intercalating agents. (See, *e.g.*, Zon, *Pharm.Res.* 5: 539-549 (1988)). To this end, the oligonucleotide may be conjugated to another molecule (*e.g.*, a peptide, hybridization triggered cross-linking agent, transport agent, hybridization-triggered cleavage agent).

25 The antisense molecules are delivered to cells that express a PDE4D polypeptide *in vivo*. A number of methods can be used for delivering antisense DNA or RNA to cells; *e.g.*, antisense molecules can be injected directly into the tissue site, or modified antisense molecules, designed to target the desired cells (*e.g.*, antisense linked to peptides or antibodies that specifically bind receptors or antigens expressed
30 on the target cell surface) can be administered systemically. Alternatively, in a another embodiment, a recombinant DNA construct is utilized in which the antisense oligonucleotide is placed under the control of a strong promoter (*e.g.*, pol III or pol

II). The use of such a construct to transfect target cells in the patient results in the transcription of sufficient amounts of single stranded RNAs that will form complementary base pairs with the endogenous transcripts and thereby prevent translation of the mRNA. For example, a vector can be introduced *in vivo* such that it is taken up by a cell and directs the transcription of an antisense RNA. Such a vector can remain episomal or become chromosomally integrated, as long as it can be transcribed to produce the desired antisense RNA. Such vectors can be constructed by recombinant DNA technology methods standard in the art and described above. For example, a plasmid, cosmid, YAC or viral vector can be used to prepare the recombinant DNA construct that can be introduced directly into the tissue site. Alternatively, viral vectors can be used which selectively infect the desired tissue, in which case administration may be accomplished by another route (*e.g.*, systemically).

In another embodiment of the invention, small double-stranded interfering RNA (RNA interference (RNAi)) can be used. RNAi is a post-transcription process, in which double-stranded RNA is introduced, and sequence-specific gene silencing results, though catalytic degradation of the targeted mRNA. See, *e.g.*, Elbashir, S.M. *et al.*, *Nature* 411:494-498 (2001); Lee, N.S., *Nature Biotech.* 19:500-505 (2002); Lee, S-K. *et al.*, *Nature Medicine* 8(7):681-686 (2002); the entire teachings of these references are incorporated herein by reference.

Endogenous expression of a gene product can also be reduced by inactivating or "knocking out" the gene or its promoter using targeted homologous recombination (*e.g.*, see Smithies *et al.*, *Nature* 317:230-234 (1985); Thomas & Capecchi, *Cell* 51:503-512 (1987); Thompson *et al.*, *Cell* 5:313-321 (1989)). For example, an altered, non-functional gene (or a completely unrelated DNA sequence) flanked by DNA homologous to the endogenous gene (either the coding regions or regulatory regions of the gene) can be used, with or without a selectable marker and/or a negative selectable marker, to transfect cells that express the gene *in vivo*. Insertion of the DNA construct, via targeted homologous recombination, results in inactivation of the gene. The recombinant DNA constructs can be directly administered or targeted to the required site *in vivo* using appropriate vectors, as described above. Alternatively, expression of non-altered genes can be increased using a similar

method: targeted homologous recombination can be used to insert a DNA construct comprising a non-altered functional gene, or the complement thereof, or a portion thereof, in place of an gene in the cell, as described above. In another embodiment, targeted homologous recombination can be used to insert a DNA construct
5 comprising a nucleic acid that encodes a polypeptide variant that differs from that present in the cell.

Alternatively, endogenous expression of a gene product can be reduced by targeting deoxyribonucleotide sequences complementary to the regulatory region (*i.e.*, the promoter and/or enhancers) to form triple helical structures that prevent
10 transcription of the gene in target cells in the body. (See generally, Helene, C., *Anticancer Drug Des.*, 6(6):569-84 (1991); Helene, C. *et al.*, *Ann. N.Y. Acad. Sci.* 660:27-36 (1992); and Maher, L. J., *Bioassays* 14(12):807-15 (1992)). Likewise, the antisense constructs described herein, by antagonizing the normal biological activity of the gene product, can be used in the manipulation of tissue, *e.g.*, tissue
15 differentiation, both *in vivo* and *for ex vivo* tissue cultures. Furthermore, the antisense techniques (*e.g.*, microinjection of antisense molecules, or transfection with plasmids whose transcripts are anti-sense with regard to a nucleic acid RNA or nucleic acid sequence) can be used to investigate the role of one or more members of the PDE4D pathway in the development of disease-related conditions. Such
20 techniques can be utilized in cell culture, but can also be used in the creation of transgenic animals.

The therapeutic agents as described herein can be delivered in a composition, as described above, or alone. They can be administered systemically, or can be targeted to a particular tissue. The therapeutic agents can be produced by a variety of
25 means, including chemical synthesis; recombinant production; *in vivo* production (*e.g.*, a transgenic animal, such as U.S. Patent No. 4,873,316 to Meade *et al.*), for example, and can be isolated using standard means such as those described herein. In addition, a combination of any of the above methods of treatment (*e.g.*, administration of non-altered polypeptide in conjunction with antisense therapy
30 targeting altered mRNA; administration of a first splicing variant in conjunction with antisense therapy targeting a second splicing variant) can also be used.

The invention additionally pertains to use of such therapeutic agents, as described herein, for the manufacture of a medicament for the treatment of stroke, TIA, MI, and/or atherosclerosis, e.g., using the methods described herein.

5 MONITORING PROGRESS OF TREATMENT

The current invention also pertains to methods of monitoring the effectiveness of treatment on the regulation of expression (*e.g.*, relative or absolute expression) of one or more PDE4D isoforms at the RNA or protein level or its enzymatic activity. PDE4D message or protein or enzymatic activity can be
10 measured in a sample of peripheral blood or cells derived therefrom. An assessment of the levels of expression or activity can be made before and during treatment with PDE4D therapeutic agents.

For example, in one embodiment of the invention, an individual who is a member of the target population can be assessed for response to treatment with a
15 PDE4D inhibitor, by examining cAMP levels or PDE4D enzymatic activity or absolute and/or relative levels of PDE4D protein or mRNA isoforms in peripheral blood in general or specific cell subfractions or combination of cell subfractions. In addition, variation such as haplotypes or mutations within or near (within 100 to 200kb) of the PDE4D gene may be used to identify individuals who are at higher risk
20 for stroke or TIA to increase the power and efficiency of clinical trials for pharmaceutical agents to prevent or treat first or subsequent stroke. The haplotypes and other variations may be used to exclude or fractionate patients in a clinical trial who are likely to have non-cAMP or non-PDE4D pathway involvement in their stroke risk in order to enrich patients who have other pathways involved and boost
25 the power and sensitivity of the clinical trial. Such variation may be used as a pharmacogenomic test to guide selection of pharmaceutical agents for individuals.

NUCLEIC ACIDS OF THE INVENTION

Nucleic Acids, Portions and Variants

30 All nucleotide positions are relative to SEQ ID NO: 1. The nucleic acids, polypeptides and antibodies described herein can be used in methods of diagnosis of susceptibility to stroke, as well as in kits useful for diagnosis of a susceptibility to

stroke. In addition, the invention pertains to isolated nucleic acid molecules comprising a human PDE4D nucleic acid. The term, "PDE4D nucleic acid," as used herein, refers to an isolated nucleic acid molecule encoding PDE4D polypeptide. The PDE4D nucleic acid molecules of the present invention can be RNA, for example, mRNA, or DNA, such as cDNA and genomic DNA. DNA molecules can be double-stranded or single-stranded; single stranded RNA or DNA can be either the coding, or sense strand or the non-coding, or antisense strand. The nucleic acid molecule can include all or a portion of the coding sequence of the gene or nucleic acid and can further comprise additional non-coding sequences such as introns and non-coding 3' and 5' sequences (including regulatory sequences, for example, as well as promoters, transcription enhancement elements, splice donor/acceptor sites, etc.). For example, a PDE4D nucleic acid can comprise the nucleic acid of SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Tables 11 and 12, the complement thereof, or to a portion or fragment of such an isolated nucleic acid molecule (*e.g.*, cDNA or the nucleic acid) that encodes PDE4D polypeptide.

Additionally, the nucleic acid molecules of the invention can be fused to a marker sequence, for example, a sequence that encodes a polypeptide to assist in isolation or purification of the polypeptide. Such sequences include, but are not limited to, those that encode a glutathione-S-transferase (GST) fusion protein and those that encode a hemagglutinin A (HA) polypeptide marker from influenza.

An "isolated" nucleic acid molecule, as used herein, is one that is separated from nucleic acids that normally flank the gene or nucleotide sequence (as in genomic sequences) and/or has been completely or partially purified from other transcribed sequences (*e.g.*, as in an RNA library). For example, an isolated nucleic acid of the invention may be substantially isolated with respect to the complex cellular milieu in which it naturally occurs, or culture medium when produced by recombinant techniques, or chemical precursors or other chemicals when chemically synthesized. In some instances, the isolated material will form part of a composition (for example, a crude extract containing other substances), buffer system or reagent mix. In other circumstances, the material may be purified to essential homogeneity, for example as determined by PAGE or column chromatography such as HPLC.

Preferably, an isolated nucleic acid molecule comprises at least about 50, 80 or 90% (on a molar basis) of all macromolecular species present. With regard to genomic DNA, the term "isolated" also can refer to nucleic acid molecules that are separated from the chromosome with which the genomic DNA is naturally associated. For
5 example, the isolated nucleic acid molecule can contain less than about 5 kb, 4 kb, 3 kb, 2 kb, 1 kb, 0.5 kb or 0.1 kb of nucleotides which flank the nucleic acid molecule in the genomic DNA of the cell from which the nucleic acid molecule is derived.

The nucleic acid molecule can be fused to other coding or regulatory sequences and still be considered isolated. Thus, recombinant DNA contained in a
10 vector is included in the definition of "isolated" as used herein. Also, isolated nucleic acid molecules include recombinant DNA molecules in heterologous host cells, as well as partially or substantially purified DNA molecules in solution. "Isolated" nucleic acid molecules also encompass *in vivo* and *in vitro* RNA transcripts of the DNA molecules of the present invention. An isolated nucleic acid
15 molecule or nucleotide sequence can include a nucleic acid molecule or nucleotide sequence that is synthesized chemically or by recombinant means. Therefore, recombinant DNA contained in a vector is included in the definition of "isolated" as used herein. Also, isolated nucleotide sequences include recombinant DNA molecules in heterologous organisms, as well as partially or substantially purified
20 DNA molecules in solution. *In vivo* and *in vitro* RNA transcripts of the DNA molecules of the present invention are also encompassed by "isolated" nucleotide sequences. Such isolated nucleotide sequences are useful in the manufacture of the encoded polypeptide, as probes for isolating homologous sequences (*e.g.*, from other mammalian species), for gene mapping (*e.g.*, by *in situ* hybridization with
25 chromosomes), or for detecting expression of the gene in tissue (*e.g.*, human tissue), such as by Northern blot analysis.

The present invention also pertains to variant nucleic acid molecules which are not necessarily found in nature but which encode a PDE4D polypeptide (*e.g.*, a polypeptide having the amino acid sequence of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10,
30 12 or 14), or another splicing variant of PDE4D polypeptide or polymorphic variant thereof. Thus, for example, DNA molecules which comprise a sequence that is different from the naturally-occurring nucleotide sequence but which, due to the

degeneracy of the genetic code, encode a PDE4D polypeptide of the present invention are also the subject of this invention. The invention also encompasses nucleotide sequences encoding portions (fragments), or encoding variant polypeptides such as analogues or derivatives of the PDE4D polypeptide. Such variants can be naturally-occurring, such as in the case of allelic variation or single nucleotide polymorphisms, or non-naturally-occurring, such as those induced by various mutagens and mutagenic processes. Intended variations include, but are not limited to, addition, deletion and substitution of one or more nucleotides that can result in conservative or non-conservative amino acid changes, including additions and deletions. Preferably the nucleotide (and/or resultant amino acid) changes are silent or conserved; that is, they do not alter the characteristics or activity of the PDE4D polypeptide. In one embodiment, the nucleotide sequences are fragments that comprise one or more polymorphic microsatellite markers. In another embodiment, the nucleotide sequences are fragments that comprise one or more single nucleotide polymorphisms in the PDE4D gene.

Other alterations of the nucleic acid molecules of the invention can include, for example, labeling, methylation, internucleotide modifications such as uncharged linkages (*e.g.*, methyl phosphonates, phosphotriesters, phosphoamidates, carbamates), charged linkages (*e.g.*, phosphorothioates, phosphorodithioates), pendent moieties (*e.g.*, polypeptides), intercalators (*e.g.*, acridine, psoralen), chelators, alkylators, and modified linkages (*e.g.*, alpha anomeric nucleic acids). Also included are synthetic molecules that mimic nucleic acid molecules in the ability to bind to a designated sequence via hydrogen bonding and other chemical interactions. Such molecules include, for example, those in which peptide linkages substitute for phosphate linkages in the backbone of the molecule.

The invention also pertains to nucleic acid molecules that hybridize under high stringency hybridization conditions, such as for selective hybridization, to a nucleotide sequence described herein (*e.g.*, nucleic acid molecules which specifically hybridize to a nucleotide sequence encoding polypeptides described herein, and, optionally, have an activity of the polypeptide). In one embodiment, the invention includes variants described herein which hybridize under high stringency hybridization conditions (*e.g.*, for selective hybridization) to a nucleotide sequence

comprising a nucleotide sequence selected from SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Tables 11 and 12 or the complement thereof. In another embodiment, the invention includes variants described herein which hybridize under high stringency hybridization conditions (e.g., for selective hybridization) to a nucleotide sequence encoding an amino acid sequence selected from SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14 or polymorphic variant thereof. In another embodiment, the protein product of the variant that hybridizes under high stringency conditions has an activity of PDE4D.

Such nucleic acid molecules can be detected and/or isolated by specific hybridization (e.g., under high stringency conditions). "Specific hybridization," as used herein, refers to the ability of a first nucleic acid to hybridize to a second nucleic acid in a manner such that the first nucleic acid does not hybridize to any nucleic acid other than to the second nucleic acid (e.g., when the first nucleic acid has a higher similarity to the second nucleic acid than to any other nucleic acid in a sample wherein the hybridization is to be performed). "Stringency conditions" for hybridization is a term of art which refers to the incubation and wash conditions, e.g., conditions of temperature and buffer concentration, which permit hybridization of a particular nucleic acid to a second nucleic acid; the first nucleic acid may be perfectly (i.e., 100%) complementary to the second, or the first and second may share some degree of complementarity which is less than perfect (e.g., 70%, 75%, 85%, 95%). For example, certain high stringency conditions can be used which distinguish perfectly complementary nucleic acids from those of less complementarity. "High stringency conditions", "moderate stringency conditions" and "low stringency conditions" for nucleic acid hybridizations are explained on pages 2.10.1-2.10.16 and pages 6.3.1-6.3.6 in *Current Protocols in Molecular Biology* (Ausubel, F.M. et al., "Current Protocols in Molecular Biology", John Wiley & Sons, (1998), the entire teachings of which are incorporated by reference herein). The exact conditions which determine the stringency of hybridization depend not only on ionic strength (e.g., 0.2XSSC, 0.1XSSC), temperature (e.g., room temperature, 42°C, 68°C) and the concentration of destabilizing agents such as formamide or denaturing agents such as SDS, but also on factors such as the length of the nucleic acid sequence, base composition, percent mismatch between

hybridizing sequences and the frequency of occurrence of subsets of that sequence within other non-identical sequences. Thus, equivalent conditions can be determined by varying one or more of these parameters while maintaining a similar degree of identity or similarity between the two nucleic acid molecules. Typically, conditions are used such that sequences at least about 60%, at least about 70%, at least about 80%, at least about 90% or at least about 95% or more identical to each other remain hybridized to one another. By varying hybridization conditions from a level of stringency at which no hybridization occurs to a level at which hybridization is first observed, conditions which will allow a given sequence to hybridize (*e.g.*, selectively) with the most similar sequences in the sample can be determined.

Exemplary conditions are described in Krause, M.H. and S.A. Aaronson, *Methods in Enzymology*, 200:546-556 (1991). Also, in, Ausubel, *et al.*, "Current Protocols in Molecular Biology", John Wiley & Sons, (1998), which describes the determination of washing conditions for moderate or low stringency conditions. Washing is the step in which conditions are usually set so as to determine a minimum level of complementarity of the hybrids. Generally, starting from the lowest temperature at which only homologous hybridization occurs, each °C by which the final wash temperature is reduced (holding SSC concentration constant) allows an increase by 1% in the maximum extent of mismatching among the sequences that hybridize. Generally, doubling the concentration of SSC results in an increase in T_m of ~17°C. Using these guidelines, the washing temperature can be determined empirically for high, moderate or low stringency, depending on the level of mismatch sought.

For example, a low stringency wash can comprise washing in a solution containing 0.2XSSC/0.1% SDS for 10 min at room temperature; a moderate stringency wash can comprise washing in a prewarmed solution (42°C) solution containing 0.2XSSC/0.1% SDS for 15 min at 42°C; and a high stringency wash can comprise washing in prewarmed (68°C) solution containing 0.1XSSC/0.1%SDS for 15 min at 68°C. Furthermore, washes can be performed repeatedly or sequentially to obtain a desired result as known in the art. Equivalent conditions can be determined by varying one or more of the parameters given as an example, as known in the art,

while maintaining a similar degree of identity or similarity between the target nucleic acid molecule and the primer or probe used.

The percent homology or identity of two nucleotide or amino acid sequences can be determined by aligning the sequences for optimal comparison purposes (*e.g.*, gaps can be introduced in the sequence of a first sequence for optimal alignment). The nucleotides or amino acids at corresponding positions are then compared, and the percent identity between the two sequences is a function of the number of identical positions shared by the sequences (*i.e.*, % identity = # of identical positions/total # of positions x 100). When a position in one sequence is occupied by the same nucleotide or amino acid residue as the corresponding position in the other sequence, then the molecules are homologous at that position. As used herein, nucleic acid or amino acid "homology" is equivalent to nucleic acid or amino acid "identity". In certain embodiments, the length of a sequence aligned for comparison purposes is at least 30%, for example, at least 40%, in certain embodiments at least 60%, and in other embodiments at least 70%, 80%, 90% or 95% of the length of the reference sequence. The actual comparison of the two sequences can be accomplished by well-known methods, for example, using a mathematical algorithm. One, non-limiting example of such a mathematical algorithm is described in Karlin *et al.*, *Proc. Natl. Acad. Sci. USA* 90:5873-5877 (1993). Such an algorithm is incorporated into the NBLAST and XBLAST programs (version 2.0) as described in Altschul *et al.*, *Nucleic Acids Res.* 25:389-3402 (1997). When utilizing BLAST and Gapped BLAST programs, the default parameters of the respective programs (*e.g.*, NBLAST) can be used. In one embodiment, parameters for sequence comparison can be set at score=100, wordlength=12, or can be varied (*e.g.*, W=5 or W=20).

Another preferred non-limiting example of a mathematical algorithm utilized for the comparison of sequences is the algorithm of Myers and Miller, CABIOS (1989). Such an algorithm is incorporated into the ALIGN program (version 2.0) which is part of the GCG sequence alignment software package. When utilizing the ALIGN program for comparing amino acid sequences, a PAM120 weight residue table, a gap length penalty of 12, and a gap penalty of 4 can be used. Additional algorithms for sequence analysis are known in the art and include ADVANCE and

ADAM as described in Torellis and Robotti (1994) *Comput. Appl. Biosci.*, 10:3-5; and FASTA described in Pearson and Lipman (1988) *PNAS*, 85:2444-8.

In another embodiment, the percent identity between two amino acid sequences can be accomplished using the GAP program in the GCG software package (Accelrys, Cambridge, UK) using either a Blossom 63 matrix or a PAM250 matrix, and a gap weight of 12, 10, 8, 6, or 4 and a length weight of 2, 3, or 4. In yet another embodiment, the percent identity between two nucleic acid sequences can be accomplished using the GAP program in the GCG software package, using a gap weight of 50 and a length weight of 3.

10 The present invention also provides isolated nucleic acid molecules that contain a fragment or portion that hybridizes under highly stringent conditions to a nucleotide sequence comprising a nucleotide sequence selected from SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Tables 11 and 12 and the complement thereof, and also provides isolated nucleic acid
15 molecules that contain a fragment or portion that hybridizes under highly stringent conditions to a nucleotide sequence encoding an amino acid sequence selected from SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or polymorphic variant thereof. The nucleic acid fragments of the invention are at least about 15, preferably at least about 18, 20, 23 or 25 nucleotides, and can be 30, 40, 50, 100, 200 or more nucleotides in
20 length. Longer fragments, for example, 30 or more nucleotides in length, which encode antigenic polypeptides described herein are particularly useful, such as for the generation of antibodies as described below.

Probes and Primers

25 In a related aspect, the nucleic acid fragments of the invention are used as probes or primers in assays such as those described herein. "Probes" or "primers" are oligonucleotides that hybridize in a base-specific manner to a complementary strand of nucleic acid molecules. By "base specific manner" is meant that the two sequences must have a degree of nucleotide complementarity sufficient for the
30 primer or probe to hybridize. Accordingly, the primer or probe sequence is not required to be perfectly complementary to the sequence of the template. Non-complementary bases or modified bases can be interspersed into the primer or probe,

provided that base substitutions do not inhibit hybridization. The nucleic acid template may also include "non-specific priming sequences" or "nonspecific sequences" to which the primer or probe has varying degrees of complementarities. Such probes and primers include polypeptide nucleic acids, as described in Nielsen *et al.*, *Science*, 254, 1497-1500 (1991).

A probe or primer comprises a region of nucleic acid that hybridizes to at least about 15, for example about 20-25, and in certain embodiments about 40, 50 or 75, consecutive nucleotides of a nucleic acid of the invention, such as a nucleic acid comprising a contiguous nucleic acid sequence of SEQ ID NO: 1 or the complement of SEQ ID NO: 1, or a nucleic acid sequence encoding an amino acid sequence of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or polymorphic variant thereof. In certain embodiments, a probe or primer comprises 100 or fewer nucleotides, in certain embodiments, from 6 to 50 nucleotides, for example, from 12 to 30 nucleotides. In other embodiments, the probe or primer is at least 70% identical to the contiguous nucleic acid sequence or to the complement of the contiguous nucleotide sequence, for example, at least 80% identical, in certain embodiments at least 90% identical, and in other embodiments at least 95% identical, or even capable of selectively hybridizing to the contiguous nucleic acid sequence or to the complement of the contiguous nucleotide sequence. Often, the probe or primer further comprises a label, *e.g.*, radioisotope, fluorescent compound, enzyme, or enzyme co-factor.

The nucleic acid molecules of the invention such as those described above can be identified and isolated using standard molecular biology techniques and the sequence information provided herein. For example, nucleic acid molecules can be amplified and isolated by the polymerase chain reaction using synthetic oligonucleotide primers designed based on one or more of the sequences provided in SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12, and/or the complement thereof, or designed based on nucleotides based on sequences encoding one or more of the amino acid sequences provided herein. See generally *PCR Technology: Principles and Applications for DNA Amplification* (ed. H.A. Erlich, Freeman Press, NY, NY, 1992); *PCR Protocols: A Guide to Methods and Applications* (Eds. Innis, *et al.*, Academic Press, San Diego,

CA, 1990); Mattila *et al.*, *Nucleic Acids Res.*, 19:4967 (1991); Eckert *et al.*, *PCR Methods and Applications*, 1:17 (1991); PCR (eds. McPherson *et al.*, IRL Press, Oxford); and U.S. Patent 4,683,202. The nucleic acid molecules can be amplified using cDNA, mRNA or genomic DNA as a template, cloned into an appropriate
5 vector and characterized by DNA sequence analysis.

Other suitable amplification methods include the ligase chain reaction (LCR) (see Wu and Wallace, *Genomics*, 4:560 (1989), Landegren *et al.*, *Science*, 241:1077 (1988), transcription amplification (Kwoh *et al.*, *Proc. Natl. Acad. Sci. USA*, 86:1173 (1989)), and self-sustained sequence replication (Guatelli *et al.*, *Proc. Nat. Acad. Sci. USA*, 87:1874 (1990)) and nucleic acid based sequence amplification (NASBA). The
10 latter two amplification methods involve isothermal reactions based on isothermal transcription, which produce both single stranded RNA (ssRNA) and double stranded DNA (dsDNA) as the amplification products in a ratio of about 30 or 100 to 1, respectively.

15 The amplified DNA can be labeled (*e.g.*, with radiolabel or other reporter molecule) and used as a probe for screening a cDNA library derived from human cells, mRNA in zap express, ZIPLOX or other suitable vector. Corresponding clones can be isolated, DNA can obtained following *in vivo* excision, and the cloned insert can be sequenced in either or both orientations by art recognized methods to identify
20 the correct reading frame encoding a polypeptide of the appropriate molecular weight. For example, the direct analysis of the nucleotide sequence of nucleic acid molecules of the present invention can be accomplished using well-known methods that are commercially available. See, for example, Sambrook *et al.*, *Molecular Cloning, A Laboratory Manual* (2nd Ed., CSHP, New York 1989); Zyskind *et al.*,
25 *Recombinant DNA Laboratory Manual*, (Acad. Press, 1988)). Using these or similar methods, the polypeptide and the DNA encoding the polypeptide can be isolated, sequenced and further characterized.

Antisense nucleic acid molecules of the invention can be designed using the nucleotide sequences of SEQ ID NO: 1 and/or the complement of SEQ ID NO: 1,
30 and/or a portion of SEQ ID NO: 1 or the complement of SEQ ID NO: 1 and/or a sequence encoding the amino acid sequences or SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 and/or 14, or encoding a portion of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 and/or

14, (wherein any one of these may optionally comprise at least one polymorphism as shown in Tables 11 and 12) and constructed using chemical synthesis and enzymatic ligation reactions using procedures known in the art. For example, an antisense nucleic acid molecule (*e.g.*, an antisense oligonucleotide) can be chemically
5 synthesized using naturally occurring nucleotides or variously modified nucleotides designed to increase the biological stability of the molecules or to increase the physical stability of the duplex formed between the antisense and sense nucleic acids, *e.g.*, phosphorothioate derivatives and acridine substituted nucleotides can be used. Alternatively, the antisense nucleic acid molecule can be produced biologically using
10 an expression vector into which a nucleic acid molecule has been subcloned in an antisense orientation (*i.e.*, RNA transcribed from the inserted nucleic acid molecule will be of an antisense orientation to a target nucleic acid of interest).

In general, the isolated nucleic acid sequences of the invention can be used as molecular weight markers on Southern gels, and as chromosome markers that are
15 labeled to map related gene positions. The nucleic acid sequences can also be used to compare with endogenous DNA sequences in patients to identify genetic disorders (*e.g.*, a predisposition for or susceptibility to stroke), and as probes, such as to hybridize and discover related DNA sequences or to subtract out known sequences from a sample. The nucleic acid sequences can further be used to derive primers for
20 genetic fingerprinting, to raise anti-polypeptide antibodies using DNA immunization techniques, and as an antigen to raise anti-DNA antibodies or elicit immune responses. Portions or fragments of the nucleotide sequences identified herein (and the corresponding complete gene sequences) can be used in numerous ways as polynucleotide reagents. For example, these sequences can be used to: (i) map their
25 respective genes on a chromosome; and, thus, locate gene regions associated with genetic disease; (ii) identify an individual from a minute biological sample (tissue typing); and (iii) aid in forensic identification of a biological sample. Additionally, the nucleotide sequences of the invention can be used to identify and express recombinant polypeptides for analysis, characterization or therapeutic use, or as
30 markers for tissues in which the corresponding polypeptide is expressed, either constitutively, during tissue differentiation, or in diseased states. The nucleic acid sequences can additionally be used as reagents in the screening and/or diagnostic

assays described herein, and can also be included as components of kits (*e.g.*, reagent kits) for use in the screening and/or diagnostic assays described herein.

Vectors

5 Another aspect of the invention pertains to nucleic acid constructs containing a nucleic acid molecule selected from the group consisting of SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12 and the complement thereof (or a portion thereof). Yet another aspect of the invention pertains to nucleic acid constructs containing a nucleic acid molecule encoding the
10 amino acid sequence of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14 or polymorphic variant thereof. The constructs comprise a vector (*e.g.*, an expression vector) into which a sequence of the invention has been inserted in a sense or antisense orientation. As used herein, the term "vector" refers to a nucleic acid molecule capable of transporting another nucleic acid to which it has been linked.
15 One type of vector is a "plasmid", which refers to a circular double stranded DNA loop into which additional DNA segments can be ligated. Another type of vector is a viral vector, wherein additional DNA segments can be ligated into the viral genome. Certain vectors are capable of autonomous replication in a host cell into which they are introduced (*e.g.*, bacterial vectors having a bacterial origin of replication and
20 episomal mammalian vectors). Other vectors (*e.g.*, non-episomal mammalian vectors) are integrated into the genome of a host cell upon introduction into the host cell, and thereby are replicated along with the host genome. Moreover, certain vectors, expression vectors, are capable of directing the expression of genes to which they are operably linked. In general, expression vectors of utility in recombinant
25 DNA techniques are often in the form of plasmids. However, the invention is intended to include such other forms of expression vectors, such as viral vectors (*e.g.*, replication defective retroviruses, adenoviruses and adeno-associated viruses) that serve equivalent functions.

 Preferred recombinant expression vectors of the invention comprise a nucleic
30 acid molecule of the invention in a form suitable for expression of the nucleic acid molecule in a host cell. This means that the recombinant expression vectors include one or more regulatory sequences, selected on the basis of the host cells to be used

for expression, which is operably linked to the nucleic acid sequence to be expressed. Within a recombinant expression vector, "operably or operatively linked" is intended to mean that the nucleotide sequence of interest is linked to the regulatory sequence(s) in a manner which allows for expression of the nucleotide sequence (e.g., in an *in vitro* transcription/translation system or in a host cell when the vector is introduced into the host cell). The term "regulatory sequence" is intended to include promoters, enhancers and other expression control elements (e.g., polyadenylation signals). Such regulatory sequences are described, for example, in Goeddel, *Gene Expression Technology: Methods in Enzymology 185*, Academic Press, San Diego, CA (1990). Regulatory sequences include those which direct constitutive expression of a nucleotide sequence in many types of host cell and those which direct expression of the nucleotide sequence only in certain host cells (e.g., tissue-specific regulatory sequences). It will be appreciated by those skilled in the art that the design of the expression vector can depend on such factors as the choice of the host cell to be transformed and the level of expression of polypeptide desired. The expression vectors of the invention can be introduced into host cells to thereby produce polypeptides, including fusion polypeptides, encoded by nucleic acid molecules as described herein.

The recombinant expression vectors of the invention can be designed for expression of a polypeptide of the invention in prokaryotic or eukaryotic cells, e.g., bacterial cells such as *E. coli*, insect cells (using baculovirus expression vectors), yeast cells or mammalian cells. Suitable host cells are discussed further in Goeddel, *supra*. Alternatively, the recombinant expression vector can be transcribed and translated *in vitro*, for example using T7 promoter regulatory sequences and T7 polymerase.

Another aspect of the invention pertains to host cells into which a recombinant expression vector of the invention has been introduced. The terms "host cell" and "recombinant host cell" are used interchangeably herein. It is understood that such terms refer not only to the particular subject cell but also to the progeny or potential progeny of such a cell. Because certain modifications may occur in succeeding generations due to either mutation or environmental influences, such

progeny may not, in fact, be identical to the parent cell, but are still included within the scope of the term as used herein.

A host cell can be any prokaryotic or eukaryotic cell. For example, a nucleic acid molecule of the invention can be expressed in bacterial cells (*e.g.*, *E. coli*),
5 insect cells, yeast or mammalian cells (such as Chinese hamster ovary cells (CHO) or COS cells). Other suitable host cells are known to those skilled in the art.

Vector DNA can be introduced into prokaryotic or eukaryotic cells via conventional transformation or transfection techniques. As used herein, the terms "transformation" and "transfection" are intended to refer to a variety of art-
10 recognized techniques for introducing a foreign nucleic acid molecule (*e.g.*, DNA) into a host cell, including calcium phosphate or calcium chloride co-precipitation, DEAE-dextran-mediated transfection, lipofection, or electroporation. Suitable methods for transforming or transfecting host cells can be found in Sambrook, *et al.*, (*supra*), and other laboratory manuals.

15 For stable transfection of mammalian cells, it is known that, depending upon the expression vector and transfection technique used, only a small fraction of cells may integrate the foreign DNA into their genome. In order to identify and select these integrants, a gene that encodes a selectable marker (*e.g.*, for resistance to antibiotics) is generally introduced into the host cells along with the gene of interest.
20 Preferred selectable markers include those that confer resistance to drugs, such as G418, hygromycin and methotrexate. Nucleic acid molecules encoding a selectable marker can be introduced into a host cell on the same vector as the nucleic acid molecule of the invention or can be introduced on a separate vector. Cells stably transfected with the introduced nucleic acid molecule can be identified by drug
25 selection (*e.g.*, cells that have incorporated the selectable marker gene will survive, while the other cells die).

A host cell of the invention, such as a prokaryotic or eukaryotic host cell in culture, can be used to produce (*i.e.*, express) a polypeptide of the invention. Accordingly, the invention further provides methods for producing a polypeptide
30 using the host cells of the invention. In one embodiment, the method comprises culturing the host cell of invention (into which a recombinant expression vector encoding a polypeptide of the invention has been introduced) in a suitable medium

such that the polypeptide is produced. In another embodiment, the method further comprises isolating the polypeptide from the medium or the host cell.

The host cells of the invention can also be used to produce nonhuman transgenic animals. For example, in one embodiment, a host cell of the invention is a
5 fertilized oocyte or an embryonic stem cell into which a nucleic acid molecule of the invention has been introduced (e.g., an exogenous PDE4D gene, or an exogenous nucleic acid encoding PDE4D polypeptide). Such host cells can then be used to create non-human transgenic animals in which exogenous nucleotide sequences have been introduced into the genome or homologous recombinant animals in which
10 endogenous nucleotide sequences have been altered. Such animals are useful for studying the function and/or activity of the nucleotide sequence and polypeptide encoded by the sequence and for identifying and/or evaluating modulators of their activity. As used herein, a "transgenic animal" is a non-human animal, preferably a mammal, more preferably a rodent such as a rat or mouse, in which one or more of
15 the cells of the animal include a transgene. Other examples of transgenic animals include non-human primates, sheep, dogs, cows, goats, chickens and amphibians. A transgene is exogenous DNA which is integrated into the genome of a cell from which a transgenic animal develops and which remains in the genome of the mature animal, thereby directing the expression of an encoded gene product in one or more
20 cell types or tissues of the transgenic animal. As used herein, an "homologous recombinant animal" is a non-human animal, preferably a mammal, more preferably a mouse, in which an endogenous gene has been altered by homologous recombination between the endogenous gene and an exogenous DNA molecule introduced into a cell of the animal, e.g., an embryonic cell of the animal, prior to
25 development of the animal.

Methods for generating transgenic animals via embryo manipulation and microinjection, particularly animals such as mice, have become conventional in the art and are described, for example, in U.S. Patent Nos. 4,736,866 and 4,870,009, U.S. Patent No. 4,873,191 and in Hogan, *Manipulating the Mouse Embryo* (Cold Spring
30 Harbor Laboratory Press, Cold Spring Harbor, N.Y., 1986). Methods for constructing homologous recombination vectors and homologous recombinant animals are described further in Bradley (1991) *Current Opinion in Bio/Technology*,

2:823-829 and in PCT Publication Nos. WO 90/11354, WO 91/01140, WO 92/0968, and WO 93/04169. Clones of the non-human transgenic animals described herein can also be produced according to the methods described in Wilmut *et al.* (1997) *Nature*, 385:810-813 and PCT Publication Nos. WO 97/07668 and WO 97/07669.

5

POLYPEPTIDES OF THE INVENTION

The present invention also pertains to isolated polypeptides encoded by PDE4D ("PDE4D polypeptides") and fragments and variants thereof, as well as polypeptides encoded by nucleotide sequences described herein (*e.g.*, other splicing variants). The term "polypeptide" refers to a polymer of amino acids, and not to a specific length; thus, peptides, oligopeptides and proteins are included within the definition of a polypeptide. As used herein, a polypeptide is said to be "isolated" or "purified" when it is substantially free of cellular material when it is isolated from recombinant and non-recombinant cells, or free of chemical precursors or other chemicals when it is chemically synthesized. A polypeptide, however, can be joined to another polypeptide with which it is not normally associated in a cell (*e.g.*, in a "fusion protein") and still be "isolated" or "purified."

The polypeptides of the invention can be purified to homogeneity. It is understood, however, that preparations in which the polypeptide is not purified to homogeneity are useful. The critical feature is that the preparation allows for the desired function of the polypeptide, even in the presence of considerable amounts of other components. Thus, the invention encompasses various degrees of purity. In one embodiment, the language "substantially free of cellular material" includes preparations of the polypeptide having less than about 30% (by dry weight) other proteins (*i.e.*, contaminating protein), less than about 20% other proteins, less than about 10% other proteins, or less than about 5% other proteins.

When a polypeptide is recombinantly produced, it can also be substantially free of culture medium, *i.e.*, culture medium represents less than about 20%, less than about 10%, or less than about 5% of the volume of the polypeptide preparation. The language "substantially free of chemical precursors or other chemicals" includes preparations of the polypeptide in which it is separated from chemical precursors or other chemicals that are involved in its synthesis. In one embodiment, the language

"substantially free of chemical precursors or other chemicals" includes preparations of the polypeptide having less than about 30% (by dry weight) chemical precursors or other chemicals, less than about 20% chemical precursors or other chemicals, less than about 10% chemical precursors or other chemicals, or less than about 5% chemical precursors or other chemicals.

In one embodiment, a polypeptide of the invention comprises an amino acid sequence encoded by a nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12 and complements and portions thereof, *e.g.*, SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or a portion or polymorphic variant thereof. However, the polypeptides of the invention also encompass fragment and sequence variants. Variants include a substantially homologous polypeptide encoded by the same genetic locus in an organism, *i.e.*, an allelic variant, as well as other splicing variants. Variants also encompass polypeptides derived from other genetic loci in an organism, but having substantial homology to a polypeptide encoded by a nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12 and complements and portions thereof, or having substantial homology to a polypeptide encoded by a nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of nucleotide sequences encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or polymorphic variants thereof. Variants also include polypeptides substantially homologous or identical to these polypeptides but derived from another organism, *i.e.*, an ortholog. Variants also include polypeptides that are substantially homologous or identical to these polypeptides that are produced by chemical synthesis. Variants also include polypeptides that are substantially homologous or identical to these polypeptides that are produced by recombinant methods.

As used herein, two polypeptides (or a region of the polypeptides) are substantially homologous or identical when the amino acid sequences are at least about 45-55%, in certain embodiments at least about 70-75%, and in other embodiments at least about 80-85%, and in others greater than about 90% or more homologous or identical. A substantially homologous amino acid sequence,

according to the present invention, will be encoded by a nucleic acid molecule hybridizing to SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12, or portion thereof, under stringent conditions as more particularly described above, or will be encoded by a nucleic acid molecule hybridizing to a nucleic acid sequence encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, portion thereof or polymorphic variant thereof, under stringent conditions as more particularly described thereof.

The invention also encompasses polypeptides having a lower degree of identity but having sufficient similarity so as to perform one or more of the same functions performed by a polypeptide encoded by a nucleic acid molecule of the invention. Similarity is determined by conserved amino acid substitution. Such substitutions are those that substitute a given amino acid in a polypeptide by another amino acid of like characteristics. Conservative substitutions are likely to be phenotypically silent. Typically seen as conservative substitutions are the replacements, one for another, among the aliphatic amino acids Ala, Val, Leu and Ile; interchange of the hydroxyl residues Ser and Thr, exchange of the acidic residues Asp and Glu, substitution between the amide residues Asn and Gln, exchange of the basic residues Lys and Arg and replacements among the aromatic residues Phe and Tyr. Guidance concerning which amino acid changes are likely to be phenotypically silent are found in Bowie *et al.*, *Science* 247:1306-1310 (1990).

A variant polypeptide can differ in amino acid sequence by one or more substitutions, deletions, insertions, inversions, fusions, and truncations or a combination of any of these. Further, variant polypeptides can be fully functional or can lack function in one or more activities. Fully functional variants typically contain only conservative variation or variation in non-critical residues or in non-critical regions. Functional variants can also contain substitution of similar amino acids that result in no change or an insignificant change in function. Alternatively, such substitutions may positively or negatively affect function to some degree. Non-functional variants typically contain one or more non-conservative amino acid substitutions, deletions, insertions, inversions, or truncation or a substitution, insertion, inversion, or deletion in a critical residue or critical region.

Amino acids that are essential for function can be identified by methods known in the art, such as site-directed mutagenesis or alanine-scanning mutagenesis (Cunningham *et al.*, *Science*, 244:1081-1085 (1989)). The latter procedure introduces single alanine mutations at every residue in the molecule. The resulting mutant molecules are then tested for biological activity *in vitro*, or *in vitro* proliferative activity. Sites that are critical for polypeptide activity can also be determined by structural analysis such as crystallization, nuclear magnetic resonance or photoaffinity labeling (Smith *et al.*, *J. Mol. Biol.*, 224:899-904 (1992); de Vos *et al.*, *Science*, 255:306-312 (1992)).

The invention also includes polypeptide fragments of the polypeptides of the invention. Fragments can be derived from a polypeptide encoded by a nucleic acid molecule comprising SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12 or a portion thereof and the complements thereof (*e.g.*, SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or other splicing variants). However, the invention also encompasses fragments of the variants of the polypeptides described herein. As used herein, a fragment comprises at least 6 contiguous amino acids. Useful fragments include those that retain one or more of the biological activities of the polypeptide as well as fragments that can be used as an immunogen to generate polypeptide-specific antibodies.

Biologically active fragments (peptides which are, for example, 6, 9, 12, 15, 16, 20, 30, 35, 36, 37, 38, 39, 40, 50, 100 or more amino acids in length) can comprise a domain, segment, or motif that has been identified by analysis of the polypeptide sequence using well-known methods, *e.g.*, signal peptides, extracellular domains, one or more transmembrane segments or loops, ligand binding regions, zinc finger domains, DNA binding domains, acylation sites, glycosylation sites, or phosphorylation sites.

Fragments can be discrete (not fused to other amino acids or polypeptides) or can be within a larger polypeptide. Further, several fragments can be comprised within a single larger polypeptide. In one embodiment a fragment designed for expression in a host can have heterologous pre- and pro-polypeptide regions fused to the amino terminus of the polypeptide fragment and an additional region fused to the carboxyl terminus of the fragment.

The invention thus provides chimeric or fusion polypeptides. These comprise a polypeptide of the invention operatively linked to a heterologous protein or polypeptide having an amino acid sequence not substantially homologous to the polypeptide. "Operatively linked" indicates that the polypeptide and the
5 heterologous protein are fused in-frame. The heterologous protein can be fused to the N-terminus or C-terminus of the polypeptide. In one embodiment the fusion polypeptide does not affect function of the polypeptide *per se*. For example, the fusion polypeptide can be a GST-fusion polypeptide in which the polypeptide sequences are fused to the C-terminus of the GST sequences. Other types of fusion
10 polypeptides include, but are not limited to, enzymatic fusion polypeptides, for example β -galactosidase fusions, yeast two-hybrid GAL fusions, poly-His fusions and Ig fusions. Such fusion polypeptides, particularly poly-His fusions, can facilitate the purification of recombinant polypeptide. In certain host cells (*e.g.*, mammalian host cells), expression and/or secretion of a polypeptide can be increased by using a
15 heterologous signal sequence. Therefore, in another embodiment, the fusion polypeptide contains a heterologous signal sequence at its N-terminus.

EP-A-O 464 533 discloses fusion proteins comprising various portions of immunoglobulin constant regions. The Fc is useful in therapy and diagnosis and thus results, for example, in improved pharmacokinetic properties (EP-A 0232 262). In
20 drug discovery, for example, human proteins have been fused with Fc portions for the purpose of high-throughput screening assays to identify antagonists. Bennett *et al.*, *Journal of Molecular Recognition*, 8:52-58 (1995) and Johanson *et al.*, *The Journal of Biological Chemistry*, 270,16:9459-9471 (1995). Thus, this invention also encompasses soluble fusion polypeptides containing a polypeptide of the
25 invention and various portions of the constant regions of heavy or light chains of immunoglobulins of various subclasses (IgG, IgM, IgA, IgE).

A chimeric or fusion polypeptide can be produced by standard recombinant DNA techniques. For example, DNA fragments coding for the different polypeptide sequences are ligated together in-frame in accordance with conventional techniques.
30 In another embodiment, the fusion gene can be synthesized by conventional techniques including automated DNA synthesizers. Alternatively, PCR amplification of nucleic acid fragments can be carried out using anchor primers which give rise to

complementary overhangs between two consecutive nucleic acid fragments which can subsequently be annealed and re-amplified to generate a chimeric nucleic acid sequence (see Ausubel *et al.*, *Current Protocols in Molecular Biology*, 1992).

Moreover, many expression vectors are commercially available that already encode a fusion moiety (*e.g.*, a GST protein). A nucleic acid molecule encoding a polypeptide of the invention can be cloned into such an expression vector such that the fusion moiety is linked in-frame to the polypeptide.

The isolated polypeptide can be purified from cells that naturally express it, purified from cells that have been altered to express it (recombinant), or synthesized using known protein synthesis methods. In one embodiment, the polypeptide is produced by recombinant DNA techniques. For example, a nucleic acid molecule encoding the polypeptide is cloned into an expression vector, the expression vector introduced into a host cell and the polypeptide expressed in the host cell. The polypeptide can then be isolated from the cells by an appropriate purification scheme using standard protein purification techniques.

In general, polypeptides of the present invention can be used as a molecular weight marker on SDS-PAGE gels or on molecular sieve gel filtration columns using art-recognized methods. The polypeptides of the present invention can be used to raise antibodies or to elicit an immune response. The polypeptides can also be used as a reagent, *e.g.*, a labeled reagent, in assays to quantitatively determine levels of the polypeptide or a molecule to which it binds (*e.g.*, a receptor or a ligand) in biological fluids. The polypeptides can also be used as markers for cells or tissues in which the corresponding polypeptide is preferentially expressed, either constitutively, during tissue differentiation, or in a diseased state. The polypeptides can be used to isolate a corresponding binding agent, *e.g.*, receptor or ligand, such as, for example, in an interaction trap assay, and to screen for peptide or small molecule antagonists or agonists of the binding interaction.

ANTIBODIES OF THE INVENTION

Polyclonal and/or monoclonal antibodies that specifically bind one form of the gene product but not to the other form of the gene product are also provided. Antibodies are also provided that bind a portion of either the variant or the reference

gene product that contains the polymorphic site or sites. The invention provides antibodies to the polypeptides and polypeptide fragments of the invention, *e.g.*, having an amino acid sequence encoded by SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or a portion thereof, or having an amino acid sequence encoded by a nucleic acid molecule comprising all or a portion of SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12 (*e.g.*, SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or another splicing variant or portion thereof). The term "antibody" as used herein refers to immunoglobulin molecules and immunologically active portions of immunoglobulin molecules, *i.e.*, molecules that contain an antigen binding site that specifically binds an antigen. A molecule that specifically binds to a polypeptide of the invention is a molecule that binds to that polypeptide or a fragment thereof, but does not substantially bind other molecules in a sample, *e.g.*, a biological sample, which naturally contains the polypeptide. Examples of immunologically active portions of immunoglobulin molecules include F(ab) and F(ab')₂ fragments which can be generated by treating the antibody with an enzyme such as pepsin. The invention provides polyclonal and monoclonal antibodies that bind to a polypeptide of the invention. The term "monoclonal antibody" or "monoclonal antibody composition", as used herein, refers to a population of antibody molecules that contain only one species of an antigen binding site capable of immunoreacting with a particular epitope of a polypeptide of the invention. A monoclonal antibody composition thus typically displays a single binding affinity for a particular polypeptide of the invention with which it immunoreacts.

Polyclonal antibodies can be prepared as described above by immunizing a suitable subject with a desired immunogen, *e.g.*, polypeptide of the invention or fragment thereof. The antibody titer in the immunized subject can be monitored over time by standard techniques, such as with an enzyme linked immunosorbent assay (ELISA) using immobilized polypeptide. If desired, the antibody molecules directed against the polypeptide can be isolated from the mammal (*e.g.*, from the blood) and further purified by well-known techniques, such as protein A chromatography to obtain the IgG fraction. At an appropriate time after immunization, *e.g.*, when the antibody titers are highest, antibody-producing cells can be obtained from the subject

and used to prepare monoclonal antibodies by standard techniques, such as the hybridoma technique originally described by Kohler and Milstein (1975) *Nature*, 256:495-497, the human B cell hybridoma technique (Kozbor *et al.* (1983) *Immunol. Today*, 4:72), the EBV-hybridoma technique (Cole *et al.* (1985), *Monoclonal Antibodies and Cancer Therapy*, Alan R. Liss, Inc., pp. 77-96) or trioma techniques. The technology for producing hybridomas is well known (see generally *Current Protocols in Immunology* (1994) Coligan *et al.* (eds.) John Wiley & Sons, Inc., New York, NY). Briefly, an immortal cell line (typically a myeloma) is fused to lymphocytes (typically splenocytes) from a mammal immunized with an immunogen as described above, and the culture supernatants of the resulting hybridoma cells are screened to identify a hybridoma producing a monoclonal antibody that binds a polypeptide of the invention.

Any of the many well known protocols used for fusing lymphocytes and immortalized cell lines can be applied for the purpose of generating a monoclonal antibody to a polypeptide of the invention (see, *e.g.*, *Current Protocols in Immunology, supra*; Galfre *et al.* (1977) *Nature*, 266:55052; R.H. Kenneth, in *Monoclonal Antibodies: A New Dimension In Biological Analyses*, Plenum Publishing Corp., New York, New York (1980); and Lerner (1981) *Yale J. Biol. Med.*, 54:387-402. Moreover, the ordinarily skilled worker will appreciate that there are many variations of such methods that also would be useful.

Alternative to preparing monoclonal antibody-secreting hybridomas, a monoclonal antibody to a polypeptide of the invention can be identified and isolated by screening a recombinant combinatorial immunoglobulin library (*e.g.*, an antibody phage display library) with the polypeptide to thereby isolate immunoglobulin library members that bind the polypeptide. Kits for generating and screening phage display libraries are commercially available (*e.g.*, the Pharmacia *Recombinant Phage Antibody System*, Catalog No. 27-9400-01; and the Stratagene *SurfZAP™* Phage Display Kit, Catalog No. 240612). Additionally, examples of methods and reagents particularly amenable for use in generating and screening antibody display library can be found in, for example, U.S. Patent No. 5,223,409; PCT Publication No. WO 92/18619; PCT Publication No. WO 91/17271; PCT Publication No. WO 92/20791; PCT Publication No. WO 92/15679; PCT Publication No. WO 93/01288; PCT

Publication No. WO 92/01047; PCT Publication No. WO 92/09690; PCT Publication No. WO 90/02809; Fuchs *et al.* (1991) *Bio/Technology*, 9:1370-1372; Hay *et al.* (1992) *Hum. Antibod. Hybridomas*, 3:81-85; Huse *et al.* (1989) *Science*, 246:1275-1281; Griffiths *et al.* (1993) *EMBO J.*, 12:725-734.

5 Additionally, recombinant antibodies, such as chimeric and humanized monoclonal antibodies, comprising both human and non-human portions, which can be made using standard recombinant DNA techniques, are within the scope of the invention. Such chimeric and humanized monoclonal antibodies can be produced by recombinant DNA techniques known in the art.

10 In general, antibodies of the invention (*e.g.*, a monoclonal antibody) can be used to isolate a polypeptide of the invention by standard techniques, such as affinity chromatography or immunoprecipitation. A polypeptide-specific antibody can facilitate the purification of natural polypeptide from cells and of recombinantly produced polypeptide expressed in host cells. Moreover, an antibody specific for a
15 polypeptide of the invention can be used to detect the polypeptide (*e.g.*, in a cellular lysate, cell supernatant, or tissue sample) in order to evaluate the abundance and pattern of expression of the polypeptide. Antibodies can be used diagnostically to monitor protein levels in tissue as part of a clinical testing procedure, *e.g.*, to, for example, determine the efficacy of a given treatment regimen. Coupling the
20 antibody to a detectable substance can facilitate detection. Examples of detectable substances include various enzymes, prosthetic groups, fluorescent materials, luminescent materials, bioluminescent materials, and radioactive materials. Examples of suitable enzymes include horseradish peroxidase, alkaline phosphatase, β -galactosidase, or acetylcholinesterase; examples of suitable prosthetic group
25 complexes include streptavidin/biotin and avidin/biotin; examples of suitable fluorescent materials include umbelliferone, fluorescein, fluorescein isothiocyanate, rhodamine, dichlorotriazinylamine fluorescein, dansyl chloride or phycoerythrin; an example of a luminescent material includes luminol; examples of bioluminescent materials include luciferase, luciferin, and aequorin, and examples of suitable
30 radioactive material include ^{125}I , ^{131}I , ^{35}S or ^3H .

DIAGNOSTIC ASSAYS

The nucleic acids, probes, primers, polypeptides and antibodies described herein can be used in methods of diagnosis of stroke or diagnosis of a susceptibility to stroke or to a disease or condition associated with an stroke gene, such as PDE4D, as well as in kits useful for diagnosis of stroke or a susceptibility to stroke or to a disease or condition associated with PDE4D. In one embodiment, the kit useful for diagnosis of stroke or susceptibility to stroke, or to a disease or condition associated with PDE4D comprises primers as described herein, wherein the primers contain one or more of the SNPs identified herein. In parallel, definition of stroke risk associated with PDE4D/cAMP pathway is useful and novel to define subgroups of individuals who would be best treated by pharmaceutical agents acting on PDE4D and/ cAMP pathways (and vice versa).

In one embodiment of the invention, diagnosis of stroke or susceptibility to stroke (or diagnosis of or susceptibility to a disease or condition associated with PDE4D) is made by detecting a polymorphism in a PDE4D nucleic acid as described herein. The polymorphism can be an alteration in a PDE4D nucleic acid, such as the insertion or deletion of a single nucleotide, or of more than one nucleotide, resulting in a frame shift alteration; the change of at least one nucleotide, resulting in a change in the encoded amino acid; the change of at least one nucleotide, resulting in the generation of a premature stop codon; the deletion of several nucleotides, resulting in a deletion of one or more amino acids encoded by the nucleotides; the insertion of one or several nucleotides, such as by unequal recombination or gene conversion, resulting in an interruption of the coding sequence of the gene or nucleic acid; duplication of all or a part of the gene or nucleic acid; transposition of all or a part of the gene or nucleic acid; or rearrangement of all or a part of the gene or nucleic acid. More than one such alteration may be present in a single gene or nucleic acid. Such sequence changes cause an alteration in the polypeptide encoded by a PDE4D nucleic acid. For example, if the alteration is a frame shift alteration, the frame shift can result in a change in the encoded amino acids, and/or can result in the generation of a premature stop codon, causing generation of a truncated polypeptide. Alternatively, a polymorphism associated with a disease or condition associated with a PDE4D nucleic acid or a susceptibility to a disease or condition associated with a

PDE4D nucleic acid can be a synonymous alteration in one or more nucleotides (*i.e.*, an alteration that does not result in a change in the polypeptide encoded by a PDE4D nucleic acid). For diagnostic applications, there may be polymorphisms informative for prediction of disease risk that are in linkage disequilibrium with the functional polymorphism. Such a polymorphism may alter splicing sites, affect the stability or transport of mRNA, or otherwise affect the transcription or translation of the nucleic acid. A PDE4D nucleic acid that has any of the alteration described above is referred to herein as an "altered nucleic acid."

In a first method of diagnosing stroke or a susceptibility to stroke, hybridization methods, such as Southern analysis, Northern analysis, or *in situ* hybridizations, can be used (see *Current Protocols in Molecular Biology*, Ausubel, F. *et al.*, eds., John Wiley & Sons, including all supplements through 1999). For example, a biological sample from a test subject (a "test sample") of genomic DNA, RNA, or cDNA, is obtained from an individual suspected of having, being susceptible to or predisposed for, or carrying a defect for, a susceptibility to a disease or condition associated with a PDE4D nucleic acid (the "test individual"). The individual can be an adult, child, or fetus. The test sample can be from any source which contains genomic DNA, such as a blood sample, sample of amniotic fluid, sample of cerebrospinal fluid, or tissue sample from skin, muscle, buccal or conjunctival mucosa, placenta, gastrointestinal tract or other organs. A test sample of DNA from fetal cells or tissue can be obtained by appropriate methods, such as by amniocentesis or chorionic villus sampling. The DNA, RNA, or cDNA sample is then examined to determine whether a polymorphism in a stroke nucleic acid is present, and/or to determine which splicing variant(s) encoded by the PDE4D is present. The presence of the polymorphism or splicing variant(s) can be indicated by hybridization of the nucleic acid in the genomic DNA, RNA, or cDNA to a nucleic acid probe. A "nucleic acid probe," as used herein, can be a DNA probe or an RNA probe; the nucleic acid probe can contain at least one polymorphism in a PDE4D nucleic acid or contains a nucleic acid encoding a particular splicing variant of a PDE4D nucleic acid. The probe can be any of the nucleic acid molecules described above (*e.g.*, the nucleic acid, a fragment, a vector comprising the nucleic acid, a probe or primer, etc.).

To diagnose a susceptibility to stroke, a hybridization sample is formed by contacting the test sample containing PDE4D, with at least one nucleic acid probe. A preferred probe for detecting mRNA or genomic DNA is a labeled nucleic acid probe capable of hybridizing to mRNA or genomic DNA sequences described
5 herein. The nucleic acid probe can be, for example, a full-length nucleic acid molecule, or a portion thereof, such as an oligonucleotide of at least 15, 30, 50, 100, 250 or 500 nucleotides in length and sufficient to specifically hybridize under stringent conditions to appropriate mRNA or genomic DNA. For example, the nucleic acid probe can be all or a portion of SEQ ID NO: 1 which may optionally
10 comprise at least one polymorphism shown in Tables 11 and 12, or the complement thereof, or a portion thereof; or can be a nucleic acid encoding a portion of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14. Other suitable probes for use in the diagnostic assays of the invention are described above (see *e.g.*, probes and primers discussed under the heading, "Nucleic Acids of the Invention").

15 The hybridization sample is maintained under conditions that are sufficient to allow specific hybridization of the nucleic acid probe to PDE4D. "Specific hybridization", as used herein, indicates exact hybridization (*e.g.*, with no mismatches). Specific hybridization can be performed under high stringency conditions or moderate stringency conditions, for example, as described above. In a
20 particularly preferred embodiment, the hybridization conditions for specific hybridization are high stringency.

Specific hybridization, if present, is then detected using standard methods. If specific hybridization occurs between the nucleic acid probe and PDE4D in the test sample, then PDE4D has the polymorphism, or is the splicing variant, that is present
25 in the nucleic acid probe. More than one nucleic acid probe can also be used concurrently in this method. In one embodiment, specific hybridization of at least one of the nucleic acid probes is indicative of a polymorphism in PDE4D, or of the presence of a particular splicing variant encoding PDE4D and is therefore diagnostic for a susceptibility to stroke.

30 In Northern analysis (see Current Protocols in Molecular Biology, Ausubel, F. *et al.*, eds., John Wiley & Sons, *supra*) the hybridization methods described above are used to identify the presence of a polymorphism or a particular splicing variant,

associated with a susceptibility to stroke. For Northern analysis, a test sample of RNA is obtained from the individual by appropriate means. Specific hybridization of a nucleic acid probe, as described above, to RNA from the individual is indicative of a polymorphism in PDE4D, or of the presence of a particular splicing variant
5 encoded by PDE4D, and is therefore diagnostic for a susceptibility to stroke.

For representative examples of use of nucleic acid probes, see, for example, U.S. Patents No. 5,288,611 and 4,851,330.

Alternatively, a peptide nucleic acid (PNA) probe can be used instead of a nucleic acid probe in the hybridization methods described above. PNA is a DNA
10 mimic having a peptide-like, inorganic backbone, such as N-(2-aminoethyl)glycine units, with an organic base (A, G, C, T or U) attached to the glycine nitrogen via a methylene carbonyl linker (see, for example, Nielsen, P.E. *et al.*, *Bioconjugate Chemistry*, 1994, 5, American Chemical Society, p. 1 (1994)). The PNA probe can be designed to specifically hybridize to a gene having a polymorphism associated with a
15 susceptibility to stroke. Hybridization of the PNA probe to PDE4D is diagnostic for a susceptibility to stroke.

In another method of the invention, mutation analysis by restriction digestion can be used to detect a mutant gene, or genes containing a polymorphism(s), if the mutation or polymorphism in the gene results in the creation or elimination of a
20 restriction site. If a restriction site is not naturally created, one can be created by PCR that depends on the polymorphism and allows genotyping. A test sample containing genomic DNA is obtained from the individual. Nucleic acid amplification methods, including but not limited to Polymerase chain reaction (PCR), Transcription Mediated Amplifications (TMA), and Ligase Mediate
25 Amplification (LMA), can be used to amplify PDE4D. The digestion pattern of the relevant DNA fragment indicates the presence or absence of the mutation or polymorphism in PDE4D, and therefore indicates the presence or absence of this susceptibility to stroke. RFLP analysis is conducted as described (see Current Protocols in Molecular Biology, *supra*). Amplification techniques based upon
30 detection of sequence of interest using reverse dot blot technology (linear array or strips) can be used and are described, for example, in U.S. Patent No. 5,468,613.

Sequence analysis can also be used to detect specific polymorphisms in PDE4D. A test sample of DNA or RNA is obtained from the test individual. PCR or other appropriate methods can be used to amplify the gene, and/or its flanking sequences, if desired. The sequence of PDE4D, or a fragment of the gene, or cDNA, or fragment of the cDNA, or mRNA, or fragment of the mRNA, is determined, using standard methods. The sequence of the gene, gene fragment, cDNA, cDNA fragment, mRNA, or mRNA fragment is compared with the known nucleic acid sequence of the gene, cDNA (*e.g.*, SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12, or a nucleic acid sequence encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or a fragment thereof) or mRNA, as appropriate. In one embodiment, the presence of at least one of the polymorphisms in PDE4D indicates that the individual has a susceptibility to stroke.

Allele-specific oligonucleotides can also be used to detect the presence of a polymorphism in PDE4D, through the use of dot-blot hybridization of amplified oligonucleotides with allele-specific oligonucleotide (ASO) probes (see, for example, Saiki, R. *et al.*, (1986), *Nature (London)* 324:163-166). An "allele-specific oligonucleotide" (also referred to herein as an "allele-specific oligonucleotide probe") is an oligonucleotide of approximately 10-50 base pairs, preferably approximately 15-30 base pairs, that specifically hybridizes to PDE4D, and that contains a polymorphism associated with a susceptibility to stroke. An allele-specific oligonucleotide probe that is specific for particular polymorphisms in PDE4D can be prepared, using standard methods (see Current Protocols in Molecular Biology, *supra*). To identify polymorphisms in the gene that are associated with a susceptibility to stroke, a test sample of DNA is obtained from the individual. PCR can be used to amplify all or a fragment of PDE4D, and its flanking sequences. The DNA containing the amplified PDE4D (or fragment of the gene) is dot-blotted, using standard methods (see Current Protocols in Molecular Biology, *supra*), and the blot is contacted with the oligonucleotide probe. The presence of specific hybridization of the probe to the amplified PDE4D is then detected. Specific hybridization of an allele-specific oligonucleotide probe to DNA from the individual is indicative of a polymorphism in PDE4D, and is therefore indicative of a susceptibility to stroke.

The invention further provides allele-specific oligonucleotides that hybridize to the reference or variant allele of a nucleic acid comprising a single nucleotide polymorphism or to the complement thereof. These oligonucleotides can be probes or primers.

5 An allele-specific primer hybridizes to a site on target DNA overlapping a polymorphism and only primes amplification of an allelic form to which the primer exhibits perfect complementarity. See Gibbs, *Nucleic Acid Res.* 17, 2427-2448 (1989). This primer is used in conjunction with a second primer that hybridizes at a distal site. Amplification proceeds from the two primers, resulting in a detectable
10 product that indicates the particular allelic form is present. A control is usually performed with a second pair of primers, one of which shows a single base mismatch at the polymorphic site and the other of which exhibits perfect complementarity to a distal site. The single-base mismatch prevents amplification and no detectable product is formed. The method works best when the mismatch is included in the 3'-
15 most position of the oligonucleotide aligned with the polymorphism because this position is most destabilizing to elongation from the primer (see, *e.g.*, WO 93/22456).

With the addition of such analogs as locked nucleic acids (LNAs), the size of primers and probes can be reduced to as few as 8 bases. LNAs are a novel class of
20 bicyclic DNA analogs in which the 2' and 4' positions in the furanose ring are joined via an O-methylene (oxy-LNA), S-methylene (thio-LNA), or amino methylene (amino-LNA) moiety. Common to all of these LNA variants is an affinity toward complementary nucleic acids, which is by far the highest reported for a DNA analog. For example, particular all oxy-LNA nonamers have been shown to have melting
25 temperatures of 64°C and 74°C when in complex with complementary DNA or RNA, respectively, as opposed to 28°C for both DNA and RNA for the corresponding DNA nonamer. Substantial increases in T_m are also obtained when LNA monomers are used in combination with standard DNA or RNA monomers. For primers and probes, depending on where the LNA monomers are included (*e.g.*,
30 the 3' end, the 5' end, or in the middle), the T_m could be increased considerably.

In another embodiment, arrays of oligonucleotide probes that are complementary to target nucleic acid sequence segments from an individual, can be

used to identify polymorphisms in PDE4D. For example, in one embodiment, an oligonucleotide linear array can be used. Oligonucleotide arrays typically comprise a plurality of different oligonucleotide probes that are coupled to a surface of a substrate in different known locations. These oligonucleotide arrays, also described
5 as "Genechips.TM.," have been generally described in the art, for example, U.S. Patent No. 5,143,854 and PCT patent publication Nos. WO 90/15070 and 92/10092. These arrays can generally be produced using mechanical synthesis methods or light directed synthesis methods that incorporate a combination of photolithographic methods and solid phase oligonucleotide synthesis methods. See Fodor *et al.*,
10 *Science*, 251:767-777 (1991), Pirrung *et al.*, U.S. Patent No. 5,143,854 (see also PCT Application No. WO 90/15070) and Fodor *et al.*, PCT Publication No. WO 92/10092 and U.S. Patent No. 5,424,186, the entire teachings of each of which are incorporated by reference herein. Techniques for the synthesis of these arrays using mechanical synthesis methods are described in, *e.g.*, U.S. Patent No. 5,384,261, the entire
15 teachings of which are incorporated by reference herein. In another embodiment, linear arrays or microarrays can be utilized.

Once an oligonucleotide array is prepared, a nucleic acid of interest is hybridized with the array and scanned for polymorphisms. Hybridization and scanning are generally carried out by methods described herein and also in, *e.g.*,
20 Published PCT Application Nos. WO 92/10092 and WO 95/11995, and U.S. Patent No. 5,424,186, the entire teachings of which are incorporated by reference herein. In brief, a target nucleic acid sequence that includes one or more previously identified polymorphic markers is amplified by well-known amplification techniques, *e.g.*, PCR. Typically, this involves the use of primer sequences that are complementary to
25 the two strands of the target sequence both upstream and downstream from the polymorphism. Asymmetric PCR techniques may also be used. Amplified target, generally incorporating a label, is then hybridized with the array under appropriate conditions. Upon completion of hybridization and washing of the array, the array is scanned to determine the position on the array to which the target sequence
30 hybridizes. The hybridization data obtained from the scan is typically in the form of fluorescence intensities as a function of location on the array.

Although primarily described in terms of a single detection block, *e.g.*, for detection of a single polymorphism, arrays can include multiple detection blocks, and thus be capable of analyzing multiple, specific polymorphisms. In alternate arrangements, it will generally be understood that detection blocks may be grouped
5 within a single array or in multiple, separate arrays so that varying, optimal conditions may be used during the hybridization of the target to the array. For example, it may often be desirable to provide for the detection of those polymorphisms that fall within G-C rich stretches of a genomic sequence, separately from those falling in A-T rich segments. This allows for the separate optimization of
10 hybridization conditions for each situation.

Additional description of use of oligonucleotide arrays for detection of polymorphisms can be found, for example, in U.S. Patents 5,858,659 and 5,837,832, the entire teachings of which are incorporated by reference herein.

Other methods of nucleic acid analysis can be used to detect polymorphisms
15 in PDE4D or splicing variants encoding by PDE4D. Representative methods include direct manual sequencing (Church and Gilbert, (1988), *Proc. Natl. Acad. Sci. USA* 81:1991-1995; Sanger, F. *et al.* (1977) *Proc. Natl. Acad. Sci.* 74:5463-5467; Beavis *et al.*, U.S. Patent No. 5,288,644); automated fluorescent sequencing; single-stranded conformation polymorphism assays (SSCP); clamped denaturing gel electrophoresis
20 (CDGE); denaturing gradient gel electrophoresis (DGGE) (Sheffield, V.C. *et al.* (1989) *Proc. Natl. Acad. Sci. USA* 86:232-236), mobility shift analysis (Orita, M. *et al.* (1989) *Proc. Natl. Acad. Sci. USA* 86:2766-2770), restriction enzyme analysis (Flavell *et al.* (1978) *Cell* 15:25; Geever, *et al.* (1981) *Proc. Natl. Acad. Sci. USA* 78:5081); heteroduplex analysis; chemical mismatch cleavage (CMC) (Cotton *et al.*
25 (1985) *Proc. Natl. Acad. Sci. USA* 85:4397-4401); RNase protection assays (Myers, R.M. *et al.* (1985) *Science* 230:1242); use of polypeptides which recognize nucleotide mismatches, such as *E. coli* mutS protein, for example.

In one embodiment of the invention, diagnosis of a disease or condition associated with PDE4D (*e.g.*, stroke) or a susceptibility to a disease or condition
30 associated with PDE4D (*e.g.*, stroke) can also be made by expression analysis by quantitative PCR (kinetic thermal cycling). This technique utilizing TaqMan[®] or Lightcycler[®] can be used to allow the identification of polymorphisms and whether

a patient is homozygous or heterozygous. The technique can assess the presence of an alteration in the expression or composition of the polypeptide encoded by a PDE4D nucleic acid or splicing variants encoded by a PDE4D nucleic acid. Further, the expression of the variants can be quantified as physically or functionally different.

In another embodiment of the invention, diagnosis of a susceptibility to stroke can also be made by examining expression and/or composition of an PDE4D polypeptide, by a variety of methods, including enzyme linked immunosorbent assays (ELISAs), Western blots, immunoprecipitations and immunofluorescence. A test sample from an individual is assessed for the presence of an alteration in the expression and/or an alteration in composition of the polypeptide encoded by PDE4D, or for the presence of a particular variant (*e.g.*, an isoform) encoded by PDE4D. An alteration in expression of a polypeptide encoded by PDE4D can be, for example, an alteration in the quantitative polypeptide expression (*i.e.*, the amount of polypeptide produced); an alteration in the composition of a polypeptide encoded by PDE4D is an alteration in the qualitative polypeptide expression (*e.g.*, expression of a mutant PDE4D polypeptide or of a different splicing variant or isoform). In one embodiment, detecting a particular splicing variant encoded by that PDE4D, or a particular pattern of splicing variants makes diagnosis of the disease or condition associated with PDE4D or a susceptibility to a disease or condition associated with PDE4D.

Both such alterations (quantitative and qualitative) can also be present. An "alteration" in the polypeptide expression or composition, as used herein, refers to an alteration in expression or composition in a test sample, as compared with the expression or composition of polypeptide by PDE4D in a control sample. A control sample is a sample that corresponds to the test sample (*e.g.*, is from the same type of cells), and is from an individual who is not affected by stroke. An alteration in the expression or composition of the polypeptide in the test sample, as compared with the control sample, is indicative of a susceptibility to stroke. Similarly, the presence of one or more different splicing variants or isoforms in the test sample, or the presence of significantly different amounts of different splicing variants in the test sample, as compared with the control sample, is indicative of a susceptibility to

stroke. Various means of examining expression or composition of the polypeptide encoded by PDE4D can be used, including spectroscopy, colorimetry, electrophoresis, isoelectric focusing, and immunoassays (*e.g.*, David *et al.*, U.S. Patent No. 4,376,110) such as immunoblotting (see also Current Protocols in Molecular Biology, particularly chapter 10). For example, in one embodiment, an antibody capable of binding to the polypeptide (*e.g.*, as described above), preferably an antibody with a detectable label, can be used. Antibodies can be polyclonal, or more preferably, monoclonal. An intact antibody, or a fragment thereof (*e.g.*, Fab or F(ab')₂) can be used. The term "labeled", with regard to the probe or antibody, is intended to encompass direct labeling of the probe or antibody by coupling (*i.e.*, physically linking) a detectable substance to the probe or antibody, as well as indirect labeling of the probe or antibody by reactivity with another reagent that is directly labeled. Examples of indirect labeling include detection of a primary antibody using a fluorescently labeled secondary antibody and end-labeling of a DNA probe with biotin such that it can be detected with fluorescently labeled streptavidin.

Western blotting analysis, using an antibody as described above that specifically binds to a polypeptide encoded by a mutant PDE4D, or an antibody that specifically binds to a polypeptide encoded by a non-mutant gene, or an antibody that specifically binds to a particular splicing variant encoded by PDE4D, can be used to identify the presence in a test sample of a particular splicing variant or isoform, or of a polypeptide encoded by a polymorphic or mutant PDE4D, or the absence in a test sample of a particular splicing variant or isoform, or of a polypeptide encoded by a non-polymorphic or non-mutant gene. The presence of a polypeptide encoded by a polymorphic or mutant gene, or the absence of a polypeptide encoded by a non-polymorphic or non-mutant gene, is diagnostic for a susceptibility to stroke, as is the presence (or absence) of particular splicing variants encoded by the PDE4D gene.

In one embodiment of this method, the level or amount of polypeptide encoded by PDE4D in a test sample is compared with the level or amount of the polypeptide encoded by PDE4D in a control sample. A level or amount of the polypeptide in the test sample that is higher or lower than the level or amount of the polypeptide in the control sample, such that the difference is statistically significant,

is indicative of an alteration in the expression of the polypeptide encoded by PDE4D, and is diagnostic for a susceptibility to stroke. Alternatively, the composition of the polypeptide encoded by PDE4D in a test sample is compared with the composition of the polypeptide encoded by PDE4D in a control sample (*e.g.*, the presence of
5 different splicing variants). A difference in the composition of the polypeptide in the test sample, as compared with the composition of the polypeptide in the control sample, is diagnostic for a susceptibility to stroke. In another embodiment, both the level or amount and the composition of the polypeptide can be assessed in the test sample and in the control sample. A difference in the amount or level of the
10 polypeptide in the test sample, compared to the control sample; a difference in composition in the test sample, compared to the control sample; or both a difference in the amount or level, and a difference in the composition, is indicative of a susceptibility to stroke.

In another embodiment, assessment of the splicing variant or isoform(s) of a
15 polypeptide encoded by a polymorphic or mutant PDE4D, can be performed. The assessment can be performed directly (*e.g.*, by examining the polypeptide itself), or indirectly (*e.g.*, by examining the mRNA encoding the polypeptide, such as through mRNA profiling). For example, probes or primers as described herein can be used to determine which splicing variants or isoforms are encoded by PDE4D mRNA,
20 using standard methods.

The presence in a test sample of a particular splicing variant(s) or isoform(s) associated with stroke or risk of stroke, or the absence in a test sample of a particular splicing variant(s) or isoform(s) not associated with stroke or risk of stroke, is diagnostic for a disease or condition associated with a PDE4D gene or a
25 susceptibility to a disease or condition associated with a PDE4D gene. Similarly, the absence in a test sample of a particular splicing variant(s) or isoform(s) associated with stroke or risk of stroke, or the presence in a test sample of a particular splicing variant(s) or isoform(s) not associated with stroke or risk of stroke, is diagnostic for the absence of disease or condition associated with a PDE4D gene or a susceptibility
30 to a disease or condition associated with a PDE4D gene.

In another embodiment, differential expression of isoforms PDE4D7, PDE4D9 and combinations thereof can be assessed and compared to control

individuals. Decreased expression of these isoforms is indicative of susceptibility to stroke, particularly carotid stroke and/or cardiogenic stroke.

The invention further pertains to a method for the diagnosis and identification of susceptibility to stroke in an individual, by identifying an at-risk haplotype in
5 PDE4D. In one embodiment, the at-risk haplotype is a haplotype for which the presence of the haplotype increases the risk of stroke significantly. Although it is to be understood that identifying whether a risk is significant may depend on a variety of factors, including the specific disease, the haplotype, and often, environmental factors, the significance may be measured by an odds ratio or a percentage. In a
10 further embodiment, the significance is measured by a percentage. In one embodiment, a significant risk is measured as an odds ratio of at least about 1.2, including but not limited to: 1.2, 1.3, 1.4, 1.5, 1.6, 1.7, 1.8 and 1.9. In a further embodiment, an odds ratio of at least 1.2 is significant. In a further embodiment, an odds ratio of at least about 1.5 is significant. In a further embodiment, a significant
15 increase in risk is at least about 1.7 is significant. In a further embodiment, a significant increase in risk is at least about 20%, including but not limited to about 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95% and 98%. In a further embodiment, a significant increase in risk is at least about 50%. It is understood however, that identifying whether a risk is medically
20 significant may also depend on a variety of factors, including the specific disease, the haplotype, and often, environmental factors.

The invention also pertains to methods of diagnosing stroke or a susceptibility to stroke in an individual, comprising screening for an at-risk haplotype in the PDE4D nucleic acid that is more frequently present in an individual
25 susceptible to stroke (affected), compared to the frequency of its presence in a healthy individual (control), wherein the presence of the haplotype is indicative of stroke or susceptibility to stroke. Standard techniques for genotyping for the presence of SNPs and/or microsatellite markers that are associated with stroke can be used, such as fluorescent-based techniques (Chen, *et al.*, *Genome Res.* 9, 492 (1999),
30 PCR, LCR, Nested PCR and other techniques for nucleic acid amplification. In one embodiment, the method comprises assessing in an individual the presence or frequency of SNPs and/or microsatellites in the PDE4D nucleic acid that are

associated with stroke, wherein an excess or higher frequency of the SNPs and/or microsatellites compared to a healthy control individual is indicative that the individual has stroke or is susceptible to stroke.

See Table 2C, Table 3, Table 4A, and 4B for SNPs and markers that comprise
5 haplotypes that can be used as screening tools. See also, Table 5, Table 6, Table 11 and Table 12 that set forth previously known SNP and novel microsatellite markers and their counterpart sequence ID reference numbers. SNPs and markers from these lists represent at-risk haplotypes and can be used to design diagnostic tests for determining a susceptibility to stroke.

10 Kits (*e.g.*, reagent kits) useful in the methods of diagnosis comprise components useful in any of the methods described herein, including for example, hybridization probes or primers as described herein (*e.g.*, labeled probes or primers), reagents for detection of labeled molecules, restriction enzymes (*e.g.*, for RFLP analysis), allele-specific oligonucleotides, antibodies which bind to altered or to non-
15 altered (native) PDE4D polypeptide, means for amplification of nucleic acids comprising PDE4D, or means for analyzing the nucleic acid sequence of PDE4D or for analyzing the amino acid sequence of an PDE4D polypeptide, etc. In one embodiment, a kit for diagnosing susceptibility to stroke can comprise primers for nucleic acid amplification of a region in the PDE4D gene comprising an at-risk
20 haplotype that is more frequently present in an individual susceptible to stroke. The primers can be designed using portions of the nucleic acids flanking SNPs that are indicative of stroke. In a particularly preferred embodiment, the primers are designed to amplify regions of the PDE4D gene associated with an at-risk haplotype for stroke, shown in Tables 8A and 8B. In another embodiment of the invention, a
25 kit for diagnosing susceptibility to stroke can further comprise probes designed to hybridize to regions of the PDE4D gene associated with an at-risk haplotype for stroke, shown in Table 5 and table 6 and/or generated from SEQ ID Nos: 85-102.

SCREENING ASSAYS AND AGENTS IDENTIFIED THEREBY

30 The invention provides methods (also referred to herein as "screening assays") for identifying the presence of a nucleotide that hybridizes to a nucleic acid of the invention, as well as for identifying the presence of a polypeptide encoded by

a nucleic acid of the invention. In one embodiment, the presence (or absence) of a nucleic acid molecule of interest (*e.g.*, a nucleic acid that has significant homology with a nucleic acid of the invention) in a sample can be assessed by contacting the sample with a nucleic acid comprising a nucleic acid of the invention (*e.g.*, a nucleic acid having the sequence of SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12, or the complement thereof, or a nucleic acid encoding an amino acid having the sequence of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or a fragment or variant of such nucleic acids), under stringent conditions as described above, and then assessing the sample for the presence (or absence) of hybridization. In another embodiment, high stringency conditions are conditions appropriate for selective hybridization. In another embodiment, a sample containing the nucleic acid molecule of interest is contacted with a nucleic acid containing a contiguous nucleotide sequence (*e.g.*, a primer or a probe as described above) that is at least partially complementary to a part of the nucleic acid molecule of interest (*e.g.*, a PDE4D nucleic acid), and the contacted sample is assessed for the presence or absence of hybridization. In another embodiment, the nucleic acid containing a contiguous nucleotide sequence is completely complementary to a part of the nucleic acid molecule of interest.

In any of these embodiments, all or a portion of the nucleic acid of interest can be subjected to amplification prior to performing the hybridization.

In another embodiment, the presence (or absence) of a polypeptide of interest, such as a polypeptide of the invention or a fragment or variant thereof, in a sample can be assessed by contacting the sample with an antibody that specifically hybridizes to the polypeptide of interest (*e.g.*, an antibody such as those described above), and then assessing the sample for the presence (or absence) of binding of the antibody to the polypeptide of interest.

In another embodiment, the invention provides methods for identifying agents (*e.g.*, fusion proteins, polypeptides, peptidomimetics, prodrugs, receptors, binding agents, antibodies, small molecules or other drugs, or ribozymes) that alter (*e.g.*, increase or decrease) the activity of the polypeptides described herein, or which otherwise interact with the polypeptides herein. For example, such agents can be agents which bind to polypeptides described herein (*e.g.*, PDE4D binding agents);

which have a stimulatory or inhibitory effect on, for example, activity of polypeptides of the invention; or which change (*e.g.*, enhance or inhibit) the ability of the polypeptides of the invention to interact with PDE4D binding agents (*e.g.*, receptors or other binding agents); or which alter posttranslational processing of the PDE4D polypeptide (*e.g.*, agents that alter proteolytic processing to direct the polypeptide from where it is normally synthesized to another location in the cell, such as the cell surface); agents that alter proteolytic processing such that more polypeptide is released from the cell, etc.

In one embodiment, the invention provides assays for screening candidate or test agents that bind to or modulate the activity of polypeptides described herein (or biologically active portion(s) thereof), as well as agents identifiable by the assays. Test agents can be obtained using any of the numerous approaches in combinatorial library methods known in the art, including: biological libraries; spatially addressable parallel solid phase or solution phase libraries; synthetic library methods requiring deconvolution; the 'one-bead one-compound' library method; and synthetic library methods using affinity chromatography selection. The biological library approach is limited to polypeptide libraries, while the other four approaches are applicable to polypeptide, non-peptide oligomer or small molecule libraries of compounds (Lam, K.S. (1997) *Anticancer Drug Des.*, 12:145).

In one embodiment, to identify agents which alter the activity of a PDE4D polypeptide, a cell, cell lysate, or solution containing or expressing a PDE4D polypeptide (*e.g.*, SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or another splicing variant encoded by PDE4D), or a fragment or derivative thereof (as described above), can be contacted with an agent to be tested; alternatively, the polypeptide can be contacted directly with the agent to be tested. The level (amount) of PDE4D activity is assessed (*e.g.*, the level (amount) of PDE4D activity is measured, either directly or indirectly), and is compared with the level of activity in a control (*i.e.*, the level of activity of the PDE4D polypeptide or active fragment or derivative thereof in the absence of the agent to be tested). If the level of the activity in the presence of the agent differs, by an amount that is statistically significant, from the level of the activity in the absence of the agent, then the agent is an agent that alters the activity of PDE4D polypeptide. An increase in the level of PDE4D activity relative to level

of the control, indicates that the agent is an agent that enhances (is an agonist of) PDE4D activity. Similarly, a decrease in the level of PDE4D activity relative to level of the control, indicates that the agent is an agent that inhibits (is an antagonist of) PDE4D activity. In another embodiment, the level of activity of a PDE4D polypeptide or derivative or fragment thereof in the presence of the agent to be tested, is compared with a control level that has previously been established. A level of the activity in the presence of the agent that differs from the control level by an amount that is statistically significant indicates that the agent alters PDE4D activity.

The present invention also relates to an assay for identifying agents which alter the expression of the PDE4D gene (*e.g.*, antisense nucleic acids, fusion proteins, polypeptides, peptidomimetics, prodrugs, receptors, binding agents, antibodies, small molecules or other drugs, or ribozymes) which alter (*e.g.*, increase or decrease) expression (*e.g.*, transcription or translation) of the gene or which otherwise interact with the nucleic acids described herein, as well as agents identifiable by the assays.

For example, a solution containing a nucleic acid encoding PDE4D polypeptide (*e.g.*, PDE4D gene) can be contacted with an agent to be tested. The solution can comprise, for example, cells containing the nucleic acid or cell lysate containing the nucleic acid; alternatively, the solution can be another solution that comprises elements necessary for transcription/translation of the nucleic acid. Cells not suspended in solution can also be employed, if desired. The level and/or pattern of PDE4D expression (*e.g.*, the level and/or pattern of mRNA or of protein expressed, such as the level and/or pattern of different splicing variants) is assessed, and is compared with the level and/or pattern of expression in a control (*i.e.*, the level and/or pattern of the PDE4D expression in the absence of the agent to be tested). If the level and/or pattern in the presence of the agent differ, by an amount or in a manner that is statistically significant, from the level and/or pattern in the absence of the agent, then the agent is an agent that alters the expression of PDE4D.

Enhancement of PDE4D expression indicates that the agent is an agonist of PDE4D activity. Similarly, inhibition of PDE4D expression indicates that the agent is an antagonist of PDE4D activity. In another embodiment, the level and/or pattern of PDE4D polypeptide(s) (*e.g.*, different splicing variants) in the presence of the agent to be tested, is compared with a control level and/or pattern that have previously been

established. A level and/or pattern in the presence of the agent that differs from the control level and/or pattern by an amount or in a manner that is statistically significant indicates that the agent alters PDE4D expression. In one embodiment, agents that can alter expression levels of isoforms PDE4D7 and/or PDE4D9 can be assessed, preferably to complement the expression levels to approximate the ratios of a healthy individual.

In another embodiment of the invention, agents which alter the expression of the PDE4D gene or which otherwise interact with the nucleic acids described herein, can be identified using a cell, cell lysate, or solution containing a nucleic acid encoding the promoter region of the PDE4D gene operably linked to a reporter gene. After contact with an agent to be tested, the level of expression of the reporter gene (*e.g.*, the level of mRNA or of protein expressed) is assessed, and is compared with the level of expression in a control (*i.e.*, the level of the expression of the reporter gene in the absence of the agent to be tested). If the level in the presence of the agent differs, by an amount or in a manner that is statistically significant, from the level in the absence of the agent, then the agent is an agent that alters the expression of PDE4D, as indicated by its ability to alter expression of a gene that is operably linked to the PDE4D gene promoter. Enhancement of the expression of the reporter indicates that the agent is an agonist of PDE4D activity. Similarly, inhibition of the expression of the reporter indicates that the agent is an antagonist of PDE4D activity. In another embodiment, the level of expression of the reporter in the presence of the agent to be tested, is compared with a control level that has previously been established. A level in the presence of the agent that differs from the control level by an amount or in a manner that is statistically significant indicates that the agent alters PDE4D expression.

Agents which alter the amounts of different splicing variants encoded by PDE4D (*e.g.*, an agent which enhances activity of a first splicing variant, and which inhibits activity of a second splicing variant), as well as agents which are agonists of activity of a first splicing variant and antagonists of activity of a second splicing variant, can easily be identified using these methods described above.

In other embodiments of the invention, assays can be used to assess the impact of a test agent on the activity of a polypeptide in relation to a PDE4D binding

agent. For example, a cell that expresses a compound that interacts with PDE4D (herein referred to as a "PDE4D binding agent", which can be a polypeptide or other molecule that interacts with PDE4D, such as a receptor) is contacted with PDE4D in the presence of a test agent, and the ability of the test agent to alter the interaction between PDE4D and the PDE4D binding agent is determined. Alternatively, a cell lysate or a solution containing the PDE4D binding agent, can be used. An agent which binds to PDE4D or the PDE4D binding agent can alter the interaction by interfering with, or enhancing the ability of PDE4D to bind to, associate with, or otherwise interact with the PDE4D binding agent. Determining the ability of the test agent to bind to PDE4D or an PDE4D binding agent can be accomplished, for example, by coupling the test agent with a radioisotope or enzymatic label such that binding of the test agent to the polypeptide can be determined by detecting the labeled with ^{125}I , ^{35}S , ^{14}C or ^3H , either directly or indirectly, and the radioisotope detected by direct counting of radioemmission or by scintillation counting.

Alternatively, test agents can be enzymatically labeled with, for example, horseradish peroxidase, alkaline phosphatase, or luciferase, and the enzymatic label detected by determination of conversion of an appropriate substrate to product. It is also within the scope of this invention to determine the ability of a test agent to interact with the polypeptide without the labeling of any of the interactants. For example, a microphysiometer can be used to detect the interaction of a test agent with PDE4D or a PDE4D binding agent without the labeling of either the test agent, PDE4D, or the PDE4D binding agent. McConnell, H.M. *et al.* (1992) *Science*, 257:1906-1912. As used herein, a "microphysiometer" (e.g., Cytosensor™) is an analytical instrument that measures the rate at which a cell acidifies its environment using a light-addressable potentiometric sensor (LAPS). Changes in this acidification rate can be used as an indicator of the interaction between ligand and polypeptide. See the Examples Section for a discussion of known PDE4D binding partners. Thus, these receptors can be used to screen for compounds that are PDE4D receptor agonists for use in treating stroke or PDE4D receptor antagonists for studying stroke. The linkage data provided herein, for the first time, provides such connection to stroke. Drugs could be designed to regulate PDE4D receptor activation that in turn can be

used to regulate signaling pathways and transcription events of genes downstream, such as Cbfa1.

In another embodiment of the invention, assays can be used to identify polypeptides that interact with one or more PDE4D polypeptides, as described herein. For example, a yeast two-hybrid system such as that described by Fields and Song (Fields, S. and Song, O., *Nature* 340:245-246 (1989)) can be used to identify polypeptides that interact with one or more PDE4D polypeptides. In such a yeast two-hybrid system, vectors are constructed based on the flexibility of a transcription factor that has two functional domains (a DNA binding domain and a transcription activation domain). If the two domains are separated but fused to two different proteins that interact with one another, transcriptional activation can be achieved, and transcription of specific markers (*e.g.*, nutritional markers such as His and Ade, or color markers such as lacZ) can be used to identify the presence of interaction and transcriptional activation. For example, in the methods of the invention, a first vector is used which includes a nucleic acid encoding a DNA binding domain and also an PDE4D polypeptide, splicing variant, fragment or derivative thereof, and a second vector is used which includes a nucleic acid encoding a transcription activation domain and also a nucleic acid encoding a polypeptide which potentially may interact with the PDE4D polypeptide, splicing variant, or fragment or derivative thereof (*e.g.*, a PDE4D polypeptide binding agent or receptor). Incubation of yeast containing the first vector and the second vector under appropriate conditions (*e.g.*, mating conditions such as used in the Matchmaker™ System from Clontech) allows identification of colonies which express the markers of interest. These colonies can be examined to identify the polypeptide(s) that interact with the PDE4D polypeptide or fragment or derivative thereof. Such polypeptides may be useful as agents that alter the activity of expression of a PDE4D polypeptide, as described above.

In more than one embodiment of the above assay methods of the present invention, it may be desirable to immobilize either PDE4D, the PDE4D binding agent, or other components of the assay on a solid support, in order to facilitate separation of complexed from uncomplexed forms of one or both of the polypeptides, as well as to accommodate automation of the assay. Binding of a test agent to the polypeptide, or interaction of the polypeptide with a binding agent in the

presence and absence of a test agent, can be accomplished in any vessel suitable for containing the reactants. Examples of such vessels include microtitre plates, test tubes, and micro-centrifuge tubes. In one embodiment, a fusion protein (*e.g.*, a glutathione-S-transferase fusion protein) can be provided which adds a domain that
5 allows PDE4D or a PDE4D binding agent to be bound to a matrix or other solid support.

In another embodiment, modulators of expression of nucleic acid molecules of the invention are identified in a method wherein a cell, cell lysate, or solution containing a nucleic acid encoding PDE4D is contacted with a test agent and the
10 expression of appropriate mRNA or polypeptide (*e.g.*, splicing variant(s)) in the cell, cell lysate, or solution, is determined. The level of expression of appropriate mRNA or polypeptide(s) in the presence of the test agent is compared to the level of expression of mRNA or polypeptide(s) in the absence of the test agent. The test agent can then be identified as a modulator of expression based on this comparison.
15 For example, when expression of mRNA or polypeptide is greater (statistically significantly greater) in the presence of the test agent than in its absence, the test agent is identified as a stimulator or enhancer of the mRNA or polypeptide expression. Alternatively, when expression of the mRNA or polypeptide is less (statistically significantly less) in the presence of the test agent than in its absence,
20 the test agent is identified as an inhibitor of the mRNA or polypeptide expression. The level of mRNA or polypeptide expression in the cells can be determined by methods described herein for detecting mRNA or polypeptide.

This invention further pertains to novel agents identified by the above-described screening assays. Accordingly, it is within the scope of this invention to
25 further use an agent identified as described herein in an appropriate animal model. For example, an agent identified as described herein (*e.g.*, a test agent that is a modulating agent, an antisense nucleic acid molecule, a specific antibody, or a polypeptide-binding agent) can be used in an animal model to determine the efficacy, toxicity, or side effects of treatment with such an agent. Alternatively, an agent
30 identified as described herein can be used in an animal model to determine the mechanism of action of such an agent. Furthermore, this invention pertains to uses of novel agents identified by the above-described screening assays for treatments as

described herein. In addition, an agent identified as described herein can be used to alter activity of a polypeptide encoded by PDE4D, or to alter expression of PDE4D, by contacting the polypeptide or the gene (or contacting a cell comprising the polypeptide or the gene) with the agent identified as described herein.

5

PHARMACEUTICAL COMPOSITIONS

The present invention also pertains to pharmaceutical compositions comprising agents described herein, particularly nucleotides encoding the polypeptides described herein; comprising polypeptides described herein (*e.g.*, one or
10 more of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14); and/or comprising other splicing variants encoded by PDE4D; and/or an agent that alters (*e.g.*, enhances or inhibits) PDE4D gene expression or PDE4D polypeptide activity as described herein. For instance, a polypeptide, protein (*e.g.*, an PDE4D receptor), an agent that alters PDE4D gene expression, or a PDE4D binding agent or binding partner, fragment,
15 fusion protein or prodrug thereof, or a nucleotide or nucleic acid construct (vector) comprising a nucleotide of the present invention, or an agent that alters PDE4D polypeptide activity, can be formulated with a physiologically acceptable carrier or excipient to prepare a pharmaceutical composition. The carrier and composition can be sterile. The formulation should suit the mode of administration.

20 Suitable pharmaceutically acceptable carriers include but are not limited to water, salt solutions (*e.g.*, NaCl), saline, buffered saline, alcohols, glycerol, ethanol, gum arabic, vegetable oils, benzyl alcohols, polyethylene glycols, gelatin, carbohydrates such as lactose, amylose or starch, dextrose, magnesium stearate, talc, silicic acid, viscous paraffin, perfume oil, fatty acid esters, hydroxymethylcellulose,
25 polyvinyl pyrrolidone, etc., as well as combinations thereof. The pharmaceutical preparations can, if desired, be mixed with auxiliary agents, *e.g.*, lubricants, preservatives, stabilizers, wetting agents, emulsifiers, salts for influencing osmotic pressure, buffers, coloring, flavoring and/or aromatic substances and the like which do not deleteriously react with the active agents.

30 The composition, if desired, can also contain minor amounts of wetting or emulsifying agents, or pH buffering agents. The composition can be a liquid solution, suspension, emulsion, tablet, pill, capsule, sustained release formulation, or

powder. The composition can be formulated as a suppository, with traditional binders and carriers such as triglycerides. Oral formulation can include standard carriers such as pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, polyvinyl pyrrolidone, sodium saccharine, cellulose, magnesium carbonate, etc.

Methods of introduction of these compositions include, but are not limited to, intradermal, intramuscular, intraperitoneal, intraocular, intravenous, subcutaneous, topical, oral and intranasal. Other suitable methods of introduction can also include gene therapy (as described below), rechargeable or biodegradable devices, particle acceleration devices ("gene guns") and slow release polymeric devices. The pharmaceutical compositions of this invention can also be administered as part of a combinatorial therapy with other agents.

The composition can be formulated in accordance with the routine procedures as a pharmaceutical composition adapted for administration to human beings. For example, compositions for intravenous administration typically are solutions in sterile isotonic aqueous buffer. Where necessary, the composition may also include a solubilizing agent and a local anesthetic to ease pain at the site of the injection. Generally, the ingredients are supplied either separately or mixed together in unit dosage form, for example, as a dry lyophilized powder or water free concentrate in a hermetically sealed container such as an ampule or sachette indicating the quantity of active agent. Where the composition is to be administered by infusion, it can be dispensed with an infusion bottle containing sterile pharmaceutical grade water, saline or dextrose/water. Where the composition is administered by injection, an ampule of sterile water for injection or saline can be provided so that the ingredients may be mixed prior to administration.

For topical application, nonsprayable forms, viscous to semi-solid or solid forms comprising a carrier compatible with topical application and having a dynamic viscosity preferably greater than water, can be employed. Suitable formulations include but are not limited to solutions, suspensions, emulsions, creams, ointments, powders, enemas, lotions, sols, liniments, salves, aerosols, etc., which are, if desired, sterilized or mixed with auxiliary agents, *e.g.*, preservatives, stabilizers, wetting agents, buffers or salts for influencing osmotic pressure, etc. The agent may be

incorporated into a cosmetic formulation. For topical application, also suitable are sprayable aerosol preparations wherein the active ingredient, preferably in combination with a solid or liquid inert carrier material, is packaged in a squeeze bottle or in admixture with a pressurized volatile, normally gaseous propellant, *e.g.*,
5 pressurized air.

Agents described herein can be formulated as neutral or salt forms. Pharmaceutically acceptable salts include those formed with free amino groups such as those derived from hydrochloric, phosphoric, acetic, oxalic, tartaric acids, etc., and those formed with free carboxyl groups such as those derived from sodium,
10 potassium, ammonium, calcium, ferric hydroxides, isopropylamine, triethylamine, 2-ethylamino ethanol, histidine, procaine, etc.

The agents are administered in a therapeutically effective amount. The amount of agents which will be therapeutically effective in the treatment of a particular disorder or condition will depend on the nature of the disorder or
15 condition, and can be determined by standard clinical techniques. In addition, *in vitro* or *in vivo* assays may optionally be employed to help identify optimal dosage ranges. The precise dose to be employed in the formulation will also depend on the route of administration, and the seriousness of the symptoms of stroke, and should be decided according to the judgment of a practitioner and each patient's circumstances.
20 Effective doses may be extrapolated from dose-response curves derived from *in vitro* or animal model test systems.

The invention also provides a pharmaceutical pack or kit comprising one or more containers filled with one or more of the ingredients of the pharmaceutical compositions of the invention. Optionally associated with such container(s) can be a
25 notice in the form prescribed by a governmental agency regulating the manufacture, use or sale of pharmaceuticals or biological products, which notice reflects approval by the agency of manufacture, use of sale for human administration. The pack or kit can be labeled with information regarding mode of administration, sequence of drug administration (*e.g.*, separately, sequentially or concurrently), or the like. The pack
30 or kit may also include means for reminding the patient to take the therapy. The pack or kit can be a single unit dosage of the combination therapy or it can be a plurality of unit dosages. In particular, the agents can be separated, mixed together

in any combination, present in a single vial or tablet. Agents assembled in a blister pack or other dispensing means is preferred. For the purpose of this invention, unit dosage is intended to mean a dosage that is dependent on the individual pharmacodynamics of each agent and administered in FDA approved dosages in
5 standard time courses.

METHODS OF THERAPY

The present invention encompasses methods of treatment (prophylactic and/or therapeutic) for stroke or a susceptibility to stroke, such as individuals in the
10 target populations described herein particularly ischemic (*e.g.*, carotid and cardiogenic strokes) and TIA, using a PDE4D therapeutic agent. A "PDE4D therapeutic agent" is an agent that alters (*e.g.*, enhances or inhibits) PDE4D polypeptide (enzymatic activity) and/or PDE4D gene expression, as described herein (*e.g.*, a PDE4D agonist or antagonist). PDE4D therapeutic agents can alter PDE4D
15 polypeptide activity or nucleic acid expression by a variety of means, such as, for example, by providing additional PDE4D polypeptide or by upregulating the transcription or translation of the PDE4D gene; by altering posttranslational processing of the PDE4D polypeptide; by altering transcription of PDE4D splicing variants; or by interfering with PDE4D polypeptide activity (*e.g.*, by binding to a
20 PDE4D polypeptide), or by downregulating the transcription or translation of the PDE4D gene.

In particular, the invention relates to methods of treatment for stroke or susceptibility to stroke (for example, for individuals in an at-risk population such as those described herein); as well as to methods of treatment for myocardial infarction, atherosclerosis, acute coronary syndrome (*e.g.*, unstable angina, non-ST-elevation
25 myocardial infarction (NSTEMI) or ST-elevation myocardial infarction (STEMI)); for decreasing risk of a second myocardial infarction; for atherosclerosis, such as for patients requiring treatment (*e.g.*, angioplasty, stents, coronary artery bypass graft) to restore blood flow in arteries (*e.g.*, coronary arteries) and peripheral arterial
30 occlusive disease.

Representative PDE4D therapeutic agents include the following:

nucleic acids or fragments or derivatives thereof described herein, particularly nucleotides encoding the polypeptides described herein and vectors comprising such nucleic acids (*e.g.*, a gene, cDNA, and/or mRNA, double-stranded interfering RNA, a nucleic acid encoding a PDE4D polypeptide or active fragment or derivative thereof, or an oligonucleotide; for example, SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12 or a nucleic acid encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or fragments or derivatives thereof), antisense nucleic acids or small double-stranded interfering RNA;

polypeptides described herein (*e.g.*, one or more of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, and/or other splicing variants encoded by PDE4D, or fragments or derivatives thereof);

other polypeptides (*e.g.*, PDE4D receptors); PDE4D binding agents; peptidomimetics; fusion proteins or prodrugs thereof; antibodies (*e.g.*, an antibody to a mutant PDE4D polypeptide, or an antibody to a non-mutant PDE4D polypeptide, or an antibody to a particular splicing variant encoded by PDE4D, as described above); ribozymes; other small molecules;

and other agents that alter (*e.g.*, inhibit or antagonize) PDE4D gene expression or polypeptide activity, or that regulate transcription of PDE4D splicing variants (*e.g.*, agents that affect which splicing variants are expressed, or that affect the amount of each splicing variant that is expressed).

More than one PDE4D therapeutic agent can be used concurrently, if desired.

The PDE4D therapeutic agent that is a nucleic acid is used in the treatment of stroke. The term, "treatment" as used herein, refers not only to ameliorating symptoms associated with the disease, but also preventing or delaying the onset of the disease, and also lessening the severity or frequency of symptoms of the disease, preventing or delaying the occurrence of a second episode of the disease or condition; and/or also lessening the severity or frequency of symptoms of the disease or condition. In the case of atherosclerosis, "treatment" also refers to a minimization or reversal of the development of plaques. The therapy is designed to alter (*e.g.*, inhibit or enhance), replace or supplement activity of a PDE4D polypeptide in an individual. For example, a PDE4D therapeutic agent can be administered in order to

upregulate or increase the expression or availability of the PDE4D gene or of specific splicing variants of PDE4D, or, conversely, to downregulate or decrease the expression or availability of the PDE4D gene or specific splicing variants of PDE4D. Upregulation or increasing expression or availability of a native PDE4D gene or of a particular splicing variant could interfere with or compensate for the expression or activity of a defective gene or another splicing variant; downregulation or decreasing expression or availability of a native PDE4D gene or of a particular splicing variant could minimize the expression or activity of a defective gene or the particular splicing variant and thereby minimize the impact of the defective gene or the particular splicing variant.

The PDE4D therapeutic agent(s) are administered in a therapeutically effective amount (*i.e.*, an amount that is sufficient to treat the disease, such as by ameliorating symptoms associated with the disease, preventing or delaying the onset of the disease, and/or also lessening the severity or frequency of symptoms of the disease). The amount which will be therapeutically effective in the treatment of a particular individual's disorder or condition will depend on the symptoms and severity of the disease, and can be determined by standard clinical techniques. In addition, *in vitro* or *in vivo* assays may optionally be employed to help identify optimal dosage ranges. The precise dose to be employed in the formulation will also depend on the route of administration, and the seriousness of the disease or disorder, and should be decided according to the judgment of a practitioner and each patient's circumstances. Effective doses may be extrapolated from dose-response curves derived from *in vitro* or animal model test systems.

In one embodiment, a nucleic acid of the invention (*e.g.*, a nucleic acid encoding a PDE4D polypeptide, such as SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12; or another nucleic acid that encodes a PDE4D polypeptide or a splicing variant, derivative or fragment thereof, such as a nucleic acid encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14) can be used, either alone or in a pharmaceutical composition as described above. For example, PDE4D or a cDNA encoding the PDE4D polypeptide, either by itself or included within a vector, can be introduced into cells (either *in vitro* or *in vivo*) such that the cells produce native PDE4D polypeptide. If necessary, cells that have

been transformed with the gene or cDNA or a vector comprising the gene or cDNA can be introduced (or re-introduced) into an individual affected with the disease. Thus, cells which, in nature, lack native PDE4D expression and activity, or have mutant PDE4D expression and activity, or have expression of a disease-associated PDE4D splicing variant, can be engineered to express PDE4D polypeptide or an active fragment of the PDE4D polypeptide (or a different variant of PDE4D polypeptide). In another embodiment, nucleic acid encoding the PDE4D polypeptide, or an active fragment or derivative thereof, can be introduced into an expression vector, such as a viral vector, and the vector can be introduced into appropriate cells in an animal. Other gene transfer systems, including viral and nonviral transfer systems, can be used. Alternatively, nonviral gene transfer methods, such as calcium phosphate coprecipitation, mechanical techniques (*e.g.*, microinjection); membrane fusion-mediated transfer via liposomes; or direct DNA uptake, can also be used.

Alternatively, in another embodiment of the invention, a nucleic acid of the invention; a nucleic acid complementary to a nucleic acid of the invention; or a portion of such a nucleic acid (*e.g.*, an oligonucleotide as described below), can be used in "antisense" therapy, in which a nucleic acid (*e.g.*, an oligonucleotide) which specifically hybridizes to the mRNA and/or genomic DNA of PDE4D is administered or generated *in situ*. The antisense nucleic acid that specifically hybridizes to the mRNA and/or DNA inhibits expression of the PDE4D polypeptide, *e.g.*, by inhibiting translation and/or transcription. Binding of the antisense nucleic acid can be by conventional base pair complementarity, or, for example, in the case of binding to DNA duplexes, through specific interaction in the major groove of the double helix.

An antisense construct of the present invention can be delivered, for example, as an expression plasmid as described above. When the plasmid is transcribed in the cell, it produces RNA that is complementary to a portion of the mRNA and/or DNA that encodes PDE4D polypeptide. Alternatively, the antisense construct can be an oligonucleotide probe that is generated *ex vivo* and introduced into cells; it then inhibits expression by hybridizing with the mRNA and/or genomic DNA of PDE4D. In one embodiment, the oligonucleotide probes are modified oligonucleotides that

are resistant to endogenous nucleases, *e.g.*, exonucleases and/or endonucleases, thereby rendering them stable *in vivo*. Exemplary nucleic acid molecules for use as antisense oligonucleotides are phosphoramidate, phosphothioate and methylphosphonate analogs of DNA (see also U.S. Patent Nos. 5,176,996; 5,264,564; and 5,256,775). Additionally, general approaches to constructing oligomers useful in antisense therapy are also described, for example, by Van der Krol *et al.* ((1988) *Biotechniques* 6:958-976); and Stein *et al.* ((1988) *Cancer Res* 48:2659-2668). With respect to antisense DNA, oligodeoxyribonucleotides derived from the translation initiation site, *e.g.*, between the -10 and +10 regions of PDE4D sequence, are preferred.

To perform antisense therapy, oligonucleotides (mRNA, cDNA or DNA) are designed that are complementary to mRNA encoding PDE4D. The antisense oligonucleotides bind to PDE4D mRNA transcripts and prevent translation. Absolute complementarity, although preferred, is not required. a sequence "complementary" to a portion of an RNA, as referred to herein, indicates that a sequence has sufficient complementarity to be able to hybridize with the RNA, forming a stable duplex; in the case of double-stranded antisense nucleic acids, a single strand of the duplex DNA may thus be tested, or triplex formation may be assayed. The ability to hybridize will depend on both the degree of complementarity and the length of the antisense nucleic acid, as described in detail above. Generally, the longer the hybridizing nucleic acid, the more base mismatches with an RNA it may contain and still form a stable duplex (or triplex, as the case may be). One skilled in the art can ascertain a tolerable degree of mismatch by use of standard procedures.

The oligonucleotides used in antisense therapy can be DNA, RNA, or chimeric mixtures or derivatives or modified versions thereof, single-stranded or double-stranded. The oligonucleotides can be modified at the base moiety, sugar moiety, or phosphate backbone, for example, to improve stability of the molecule, hybridization, etc. The oligonucleotides can include other appended groups such as peptides (*e.g.*, for targeting host cell receptors *in vivo*), or agents facilitating transport across the cell membrane (see, *e.g.*, Letsinger *et al.* (1989) *Proc. Natl. Acad. Sci. USA* 86:6553-6556; Lemaitre *et al.*, (1987), *Proc. Natl. Acad. Sci. USA* 84:648-652; PCT International Publication No. W088/09810) or the blood-brain barrier (see, *e.g.*,

PCT International Publication No. W089/10134), or hybridization-triggered cleavage agents (see, *e.g.*, Krol *et al.* (1988) *BioTechniques* 6:958-976) or intercalating agents. (See, *e.g.*, Zon, (1988), *Pharm. Res.* 5:539-549). To this end, the oligonucleotide may be conjugated to another molecule (*e.g.*, a peptide, hybridization triggered cross-
5 linking agent, transport agent, hybridization-triggered cleavage agent).

The antisense molecules are delivered to cells that express PDE4D *in vivo*. A number of methods can be used for delivering antisense DNA or RNA to cells; *e.g.*, antisense molecules can be injected directly into the tissue site, or modified antisense molecules, designed to target the desired cells (*e.g.*, antisense linked to peptides or
10 antibodies that specifically bind receptors or antigens expressed on the target cell surface) can be administered systematically. Alternatively, in another embodiment, a recombinant DNA construct is utilized in which the antisense oligonucleotide is placed under the control of a strong promoter (*e.g.*, pol III or pol II). The use of such a construct to transfect target cells in the patient results in the transcription of
15 sufficient amounts of single stranded RNAs that will form complementary base pairs with the endogenous PDE4D transcripts and thereby prevent translation of the PDE4D mRNA. For example, a vector can be introduced *in vivo* such that it is taken up by a cell and directs the transcription of an antisense RNA. Such a vector can remain episomal or become chromosomally integrated, as long as it can be
20 transcribed to produce the desired antisense RNA. Such vectors can be constructed by recombinant DNA technology methods standard in the art and described above. For example, a plasmid, cosmid, YAC or viral vector can be used to prepare the recombinant DNA construct that can be introduced directly into the tissue site. Alternatively, viral vectors can be used which selectively infect the desired tissue, in
25 which case administration may be accomplished by another route (*e.g.*, systemically).

Methods of modulating PDE4D expression by administering an RNA inhibitor of the activity of the target protein are also possible. The term "RNA inhibitor" refers to an inhibitory RNA that silences expression of the target protein
30 by RNA interference (McManus, M.T. and Sharp, P.A., 2002. *Nat. Rev. Genet.* 3:737-47; Hannon, G.J., 2002. *Nature* 418:244-51; Paddison, P.J. and Hannon, G.J., 2002. *Cancer Cell* 2:17-23). RNA interference is conserved throughout evolution,

from *C. elegans* to humans, and is believed to function in protecting cells from invasion by RNA viruses. When a cell is infected by a dsRNA virus, the dsRNA is recognized and targeted for cleavage by an RNaseIII-type enzyme termed Dicer. The Dicer enzyme "dices" the RNA into short duplexes of 21 nucleotides, termed
5 short-interfering RNAs or siRNAs, composed of 19 nucleotides of perfectly paired ribonucleotides with two unpaired nucleotides on the 3' end of each strand. These short duplexes associate with a multiprotein complex termed RISC, and direct this complex to mRNA transcripts with sequence similarity to the siRNA. As a result, nucleases present in the RISC complex cleave the mRNA transcript, thereby
10 abolishing expression of the gene product. In the case of viral infection, this mechanism would result in destruction of viral transcripts, thus preventing viral synthesis. Since the siRNAs are double-stranded, either strand has the potential to associate with RISC and direct silencing of transcripts with sequence similarity.

Recently, it was determined that gene silencing could be induced by
15 presenting the cell with the siRNA, mimicking the product of Dicer cleavage (Elbashir, S.M., *et al.*, 2001. *Nature* 411:494-8; Elbashir, S.M., *et al.*, 2001. *Genes Dev.* 15:188-200). Synthetic siRNA duplexes maintain the ability to associate with RISC and direct silencing of mRNA transcripts, thus providing researchers with a powerful tool for gene silencing in mammalian cells. Yet another method to
20 introduce the dsRNA for gene silencing is shRNA, for short hairpin RNA (Paddison, P.J., *et al.*, 2002. *Genes Dev.* 16:948-58; Brummelkamp, T.R., *et al.*, 2002 *Science* 296:550-3; Sui, G., *et al.*, 2002. *Proc. Natl. Acad. Sci. U.S.A.* 99:5515-20). In this case, a desired siRNA sequence is expressed from a plasmid (or virus) containing an "shRNA" gene having an inverted repeat with an intervening loop sequence to form
25 a hairpin structure. The resulting shRNA transcript containing the hairpin is subsequently processed by Dicer to produce siRNAs for silencing. Plasmid-based shRNAs can be expressed stably in cells, allowing long-term gene silencing in cells, or even in animals (McCaffrey, A.P., *et al.*, 2002. *Nature* 418:38-9; Xia, H., *et al.*, 2002. *Nat. Biotech.* 20:1006-10; Lewis, D.L., *et al.*, 2002. *Nat. Genetics* 32:107-8;
30 Robinson, D.A., *et al.*, 2003. *Nat. Genetics* 33:401-6; Tiscornia, G., *et al.*, (2003) *Proc. Natl. Acad. Sci. U.S.A.* 100:1844-8). RNA interference has been successfully

used therapeutically to protect mice from fulminant hepatitis (Song, E., *et al.*, 2003. *Nat. Medicine* 9:347-51).

Endogenous PDE4D expression can be also reduced by inactivating or "knocking out" PDE4D or its promoter using targeted homologous recombination (e.g., see Smithies *et al.* (1985) *Nature* 317:230-234; Thomas & Capecchi (1987) *Cell* 51:503-512; Thompson *et al.* (1989) *Cell* 5:313-321). For example, a mutant, non-functional PDE4D (or a completely unrelated DNA sequence) flanked by DNA homologous to the endogenous PDE4D (either the coding regions or regulatory regions of PDE4D) can be used, with or without a selectable marker and/or a negative selectable marker, to transfect cells that express PDE4D *in vivo*. Insertion of the DNA construct, via targeted homologous recombination, results in inactivation of PDE4D. The recombinant DNA constructs can be directly administered or targeted to the required site *in vivo* using appropriate vectors, as described above. Alternatively, expression of non-mutant PDE4D can be increased using a similar method: targeted homologous recombination can be used to insert a DNA construct comprising a non-mutant, functional PDE4D (e.g., a gene having SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 11 and 12), or a portion thereof, in place of a mutant PDE4D in the cell, as described above. In another embodiment, targeted homologous recombination can be used to insert a DNA construct comprising a nucleic acid that encodes a PDE4D polypeptide variant that differs from that present in the cell.

Alternatively, endogenous PDE4D expression can be reduced by targeting deoxyribonucleotide sequences complementary to the regulatory region of PDE4D (*i.e.*, the PDE4D promoter and/or enhancers) to form triple helical structures that prevent transcription of PDE4D in target cells in the body. (See generally, Helene, C. (1991) *Anticancer Drug Des.*, 6(6):569-84; Helene, C., *et al.* (1992) *Ann, N.Y. Acad. Sci.*, 660:27-36; and Maher, L. J. (1992) *Bioassays* 14(12):807-15). Likewise, the antisense constructs described herein, by antagonizing the normal biological activity of one of the PDE4D proteins, can be used in the manipulation of tissue, *e.g.*, tissue differentiation, both *in vivo* and *for ex vivo* tissue cultures. Furthermore, the anti-sense techniques (*e.g.*, microinjection of antisense molecules, or transfection with plasmids whose transcripts are anti-sense with regard to a PDE4D mRNA or gene

sequence) can be used to investigate role of PDE4D in developmental events, as well as the normal cellular function of PDE4D in adult tissue. Such techniques can be utilized in cell culture, but can also be used in the creation of transgenic animals.

In yet another embodiment of the invention, other PDE4D therapeutic agents
5 as described herein can also be used in the treatment or prevention of stroke. The therapeutic agents can be delivered in a composition, as described above, or by themselves. They can be administered systemically, or can be targeted to a particular tissue. The therapeutic agents can be produced by a variety of means, including chemical synthesis; recombinant production; *in vivo* production (*e.g.*, a transgenic
10 animal, such as U.S. Patent No. 4,873,316 to Meade *et al.*), for example, and can be isolated using standard means such as those described herein.

A combination of any of the above methods of treatment (*e.g.*, administration of non-mutant PDE4D polypeptide in conjunction with antisense therapy targeting mutant PDE4D mRNA; administration of a first splicing variant encoded by PDE4D
15 in conjunction with antisense therapy targeting a second splicing encoded by PDE4D), can also be used.

The invention will be further described by the following non-limiting examples. The teachings of all publications cited herein are incorporated herein by reference in their entirety.

20

EXAMPLES

EXAMPLE 1: PDE4D VARIATIONS AND HAPLOTYPES INCREASE RISK FOR STROKE

25

Icelandic Stroke Patients and Phenotype Characterization

A population-based list containing 2543 Icelandic stroke patients, diagnosed from 1993 through 1997, was derived from two major hospitals in Iceland and the Icelandic Heart Association (the study was approved by the Icelandic Data Protection
30 Commission of Iceland and the National Bioethics Committee). Patients with hemorrhagic stroke represented 6% of all patients (patients with the Icelandic type of hereditary cerebral hemorrhage with amyloidosis and patients with subarachnoid

hemorrhage were excluded). Ischemic stroke accounted for 67% of the total patients and TIAs 27%. The distribution of stroke subtypes in this study is similar to that reported in other Caucasian populations (Mohr, J.P., *et al.*, *Neurology*, 28:754-762 (1978); L. R. Caplan, *In Stroke, A Clinical Approach* (Butterworth-Heinemann, Stoneham, MA, ed 3, (1993)).

The list of approximately 2000 living patients was run through our computerized genealogy database. A comprehensive genealogy database that has been established at deCODE genetics was used to cluster the patients in pedigrees. Each version of the computerized genealogy database was reversibly encrypted by the Data Protection Commission of Iceland before arriving at the laboratory (Gulcher, J.R., *et al.*, *Eur. J. Hum. Genet.* 8:739 (2000)). The database uses a patient list, with encrypted personal identifiers, as input, and recursive algorithms to find all ancestors in the database who are related to any member on the input list within a given number of generations back (Gulcher, J.R., and Stefansson, K., *Clin. Chem. Lab. Med.* 36:523 (1998)) covering the whole Icelandic nation. The cluster function then searches for ancestors who are common to any two or more members of the input list. One hundred and seventy-nine families with two or more living patients were chosen for the study with a total of 476 patients connected within 6 meioses (6 meioses connect second cousins). Informed consent was obtained from all patients and their relatives whose DNA samples were used in the linkage scan. The mean separation between affected pairs is 4.8 meioses. Of the patients selected for the study 73% had ischemic strokes, 23% TIAs and 4% hemorrhagic strokes.

In the selected families, hemorrhagic stroke patients clustered with ischemic stroke and TIA patients, and there were no families with a striking preponderance of hemorrhagic stroke or of the subtypes of ischemic stroke. Patients with ischemic stroke were reclassified according to the TOAST (Trial of Org 10172 in Acute Stroke Treatment) sub-classification system for stroke (Adams, H.P., Jr., *et al.*, *Stroke*, 24:34-41 (1993)). This system includes five categories: (1) large-artery atherosclerosis, (2) cardioembolism, (3) small-artery occlusion (lacune), (4) stroke of other determined etiology and (5) stroke of undetermined etiology. The diagnoses were based on clinical features and on data from ancillary diagnostic studies. Patients defined with large-artery atherosclerosis had clinical and brain imaging

findings of cerebral cortical dysfunction and either significant (>70%) stenosis (this is a stricter criteria than used in TOAST where 50% stenosis is the cut-off) or occlusion of a major brain artery or branch cortical artery. Potential sources of cardiogenic embolism were excluded. The category cardioembolism included
5 patients with at least one cardiac source for an embolus and potential large-artery sources of thrombosis and embolism was eliminated. Patients with small-artery occlusion had one of the traditional clinical lacunar syndromes and no evidence of cerebral cortical dysfunction. Potential cardiac source of embolus and stenosis >70% in an ipsilateral extracranial artery was excluded. The category, acute stroke of other
10 determined etiology, included patients with rare causes of stroke and patients with two or more potential causes of stroke. If the causes of stroke could not be determined despite extensive evaluation patients were included in the category stroke of undetermined etiology. FIG. 1 displays two pedigrees each affected by several of the stroke subtypes, including hemorrhagic stroke. Apparently what is inherited in
15 stroke is the broadly defined phenotype.

Genome-wide scan

A genome-wide scan was performed using a framework map of about 1000 microsatellite markers. The DNA samples were genotyped using approximately
20 1000 fluorescently labelled primers. A microsatellite screening set based in part on the ABI Linkage Marker (v2) screening set and the ABI Linkage Marker (v2) intercalating set in combination with 500 custom-made markers were developed. All markers were extensively tested for robustness, ease of scoring, and efficiency in 4X multiplex PCR reactions. In the framework marker set, the average spacing between
25 markers was approximately 4 cM with no gaps larger than 10 cM. Marker positions were obtained from the Marshfield map, except for a three-marker putative inversion on chromosome 8 (Jonsdottir, G.M., *et al.*, *Am. J. Hum. Genet.*, 67 (Suppl. 2):332 (2000); Yu, A., *et al.*, *Am. J. Hum. Genet.* 67 (Suppl. 2):10 (2000). The PCR amplifications were set up, run and pooled on Perkin Elmer/Applied Biosystems 877
30 Integrated Catalyst Thermocyclers with a similar protocol for each marker. The reaction volume used was 5 µl and for each PCR reaction 20 ng of genomic DNA was amplified in the presence of 2 pmol of each primer, 0.25 U AMPLITAQ GOLD

(DNA polymerase; trademark of Roche Molecular Systems), 0.2 mM dNTPs and 2.5 mM MgCl₂ (buffer was supplied by manufacturer). The PCR conditions used were 95°C for 10 minutes, then 37 cycles of 15 s at 94°C, 30s at 55°C and 1 min at 72°C. The PCR products were supplemented with the internal size standard and the pools
5 were separated and detected on Applied Biosystems model 377 Sequencer using v3.0 GENESCAN (peak calling software; trademark of Applied Biosystems). Alleles were called automatically with the TRUEALLELE (computer program for alleles identification; trademark of Cybergene, Inc.) program, and the program, DECODE-GT (computer editing program that works downstream of the
10 TRUEALLELE program; trademark of deCODE genetics), was used to fractionate according to quality and edit the called genotypes (Palsson, B., *et al.*, *Genome Res.* 9:1002 (1999)). At least 180 Icelandic controls were genotyped to derive allelic frequencies.

A total of 476 patients and 438 relatives were genotyped. The data was
15 analyzed and the statistical significance determined by applying affecteds-only allele-sharing methods (which does not specify any particular inheritance model) implemented in the ALLEGRO (computer program for multipoint linkage analysis; trademark of deCODE genetics) program that calculates lod scores based on multipoint calculations. Our baseline linkage analysis uses the S_{pairs} scoring function
20 (Kruglyak, L., *et al.*, *Am. J. Hum. Genet.*, 58:1347 (1996)), the exponential allele-sharing model (Kong, A. and Cox, N.J., *Am. J. Hum. Genet.*, 61:1179 (1997)), and a family weighting scheme which is halfway, on the log scale, between weighting each affected pair equally and weighting each family equally. In the analysis we treat all genotyped individuals who are not affected as "unknown". All linkage analyses in
25 this paper were performed using multipoint calculation with the program ALLEGRO (deCODE genetics) (Gudbjartsson, D.F., *et al.*, *Nat. Genet.* 25:12 (2000)).

The allele sharing lod scores for the genome scan using the framework map showed three regions that achieved a lod score above 1.0. Two of these regions are on chromosome 5q. The first peak is at approximately 69 cM with a lod score of
30 2.00. The second peak is at 99 cM with a lod score of 1.14. The third region is on chromosome 14q at 55 cM with a lod score of 1.24.

The information for linkage at the 5q locus was increased by genotyping an additional 45 markers over a 45 cM segment which spanned both peaks. The information used here is defined by Nicolae (D. L. Nicolae, Thesis, University of Chicago (1999)) and has been demonstrated to be asymptotically equivalent to a classical measure of the fraction of missing information (Dempster, A.P., *et al.*, *J. R. Statist. Soc. B*, 39:1 (1977)). While the lod score at the second peak dropped slightly to around 1.05, the lod score at the first peak increased to 3.39. However, close inspection of our results suggested that not only does the Marshfield genetic map lack resolution (many markers assigned the same map location), but also there may be some errors in their order. As a result, the genetic length of the region estimated using our material was substantially greater than what is reported. By modifying the ALLEGRO (deCODE genetics) program, we applied the EM algorithm to our data to estimate the genetic distances between markers. We found that our estimate of the genetic length of the region was substantially longer than that given in the Marshfield map. This indicates a problem with marker order because, in general, incorrect marker order leads to an increased number of apparent crossovers and increases the apparent genetic length.

Physical and genetic mapping

The marker order and inter-marker distances were improved by constructing high density physical and genetic maps over a 20 cM region between markers D5S474 and D5S2046. A combination of data from coincident hybridizations of BAC membranes using a high density of STSs and the Fingerprinting Contig database was used to build large contigs of BACs from the RPCI -11 library. The order of the linkage markers was also confirmed by high-resolution genetic mapping using the stroke families supplemented with over 112 other large nuclear families. High resolution genetic mapping was used both to anchor and place in order contigs found by physical mapping as well as to obtain accurate inter-marker distances for the correctly ordered markers. Data from 112 Icelandic nuclear families (sibships with their parents, containing from two to seven siblings) were analyzed together with the nuclear families available within the stroke pedigrees. For the purpose of genetic mapping the 112 nuclear families alone provide 588 meioses, and the total

number of meioses available for mapping was over 2000. By comparison, the Marshfield genetic map was constructed based on 182 meioses. The large number of meiotic events within our families provides the ability to map markers to the resolution of 0.5 to 1.0 cM. Combining this information with the physical map
5 resulted in a highly reliable order of markers and inter-marker distances within this 20 cM region. Linkage markers common to the genetic and physical maps were used to anchor and place in order four of the physically mapped contigs. By integrating the genetic and physical maps a most likely order of 30 polymorphic markers was derived.

10 BAC contigs were generated by a method that combines coincident primer hybridization with data mining. The RPCI-11 human male BAC library segments 1 & 2 (Pieter de Jong, Children's Hospital Oakland Research Institute) containing about 200,000 clones with a 12X coverage, were gridded using a 6x6 double offset pattern in 23 cm x 23 cm membranes with a BioGrid robot (Biorobotics Ltd.,
15 Cambridge, UK). Initially, hybridizations were performed with markers in the region of interest according to their location in the Weizmann Institute Unified Database. Primer sequences were analyzed and discarded according to their content of known repeats, *E. coli* and vector sequences (the analysis was performed using software developed at deCODE genetics). One hundred and fifty markers in the
20 region (30 polymorphic markers used in linkage and 120 generated from STSs) separated by an average of 130 kb were used. The selected markers were used to generate two ³²P labelled probes, F that contained the pooled forward primers and R that contained the pooled reverse primers. Reading of positive signals was performed automatically from digitized images of resulting autoradiograms by
25 informatics tools developed at deCODE genetics. The coincident signals in both hybridizations were selected as positive clones. A set of overlapping clones was assembled through a combination of hybridization and BAC fingerprint walking. Fingerprints of positive clones were analyzed using the FPC database developed at the Sanger Center. Data from FPC contigs prebuilt with a cutoff of 3e-12 and from
30 sequence datamining was integrated with the hybridization results. BACs in the region detected by data mining and hybridization were re-arrayed using a Multiprobe IIx robot (Packard, Meriden, CT). Small membranes (8 cm x 12 cm) were gridded

in 6x6 double offset pattern and individually hybridized with the markers of interest. Positive patterns were transferred using transparencies to an Excel file containing macros to provide BAC to marker associations. A visual map was generated by combining the hybridization, fingerprinting and sequence data. New markers were
5 generated from BAC end sequences to close the gap. After several rounds of hybridization positive BACs were assembled into 7 contigs covering approximately 20 Mb. Thirty of the polymorphic markers used in linkage were assigned to four of the contigs. Estimation of contig lengths and distance between markers assigned to them was based on the FPC program.

10 Twenty-seven of our 30 linkage markers mapped to three contigs in the October 2000 release from UCSC, the UC Santa Cruz (UCSC) draft assembly. The marker order within the contigs is in agreement with our order with the exception of two markers. Although the UCSC assemblies are improving, some contigs have incorrect order, orientation, or contig assembly. We believe that high resolution
15 genetic mapping and perhaps focused hybridization experiments are still necessary to confirm accuracy of sequence assemblies. In addition, high resolution genetic mapping provides better estimates of inter-marker genetic distances that are also important for linkage analysis (Halpern, J. and Whittemore, A.S., *Hum. Hered.* 49:194 (1999); Daw, E.W., *et al.*, *Genet. Epidemiol.* 19:366 (2000)).

20 *Statistical Methods for Linkage Analysis*

Multipoint, affected-only allele-sharing methods were used in the analyses to assess evidence for linkage. All results, both the LOD-score and the non-parametric linkage (NPL) score, were obtained using the program Allegro (Gudbjartsson *et al.*,
25 *Nat. Genet.* 25:12-3, 2000). Our baseline linkage analysis, as previously described (Gretarsdottir *et al.*, *Am J Hum Genet.* 70:593-603, 2002), uses the S_{pairs} scoring function (Whittemore, A.S., Halpern, J. (1994), *Biometrics* 50:118-27; Kruglyak L, *et al.* (1996), *Am J Hum Genet* 58:1347-63), the exponential allele-sharing model (Kong, A. and Cox, N.J. (1997), *Am J Hum Genet* 61:1179-88) and a family
30 weighting scheme that is halfway, on the log-scale, between weighting each affected pair equally and weighting each family equally. The information measure we use is part of the Allegro program output and the information value equals zero if the

marker genotypes are completely uninformative and equals one if the genotypes determine the exact amount of allele sharing by decent among the affected relatives (Gretarsdottir *et al.*, *Am. J. Hum. Genet.*, 70:593-603, (2002)). We computed the P-values two different ways and here report the less significant result. The first P-value
5 was computed on the basis of large sample theory; the distribution of $Z_{lr} = \sqrt{(2[\log_e(10)\text{LOD}])}$ approximates a standard normal variable under the null hypothesis of no linkage (Kong, A. and Cox, N.J. (1997), *Am J Hum Genet* 61:1179-88). The second P-value was calculated by comparing the observed LOD-score with its complete data sampling distribution under the null hypothesis (Gudbjartsson *et*
10 *al.*, *Nat. Genet.* 25:12-3, 2000). When the data consist of more than a few families, as is the case here, these two P-values tend to be very similar.

Final linkage results and localization

Linkage analysis including genotypes from the higher density markers using
15 the deCODE marker order resulted in a lod score of 4.40 ($P = 3.9 \times 10^{-6}$) on chromosome 5q12 at the marker D5S2080. The reported P value is part of the output of the ALLEGRO (deCODE genetics) program which was developed at deCODE and has become a standard linkage program worldwide over the last 3 years (Gudbjartsson *et al.*, *Nat. Genet.* 25:12-3, 2000). We have given it to over 200
20 academic departments around the world free of charge and it is widely used. The locus has been designated as *STRK1*. With the addition of these extra markers, it was possible to narrow down the region to a segment less than 6 cM, from D5S1474 to D5S398, as defined by one drop in lod.

To further investigate the contribution of this susceptibility locus to stroke, a
25 range of parametric models were fitted to the data. However, all analyses were still *affecteds only* in the sense that individuals were either classified as affecteds or having unknown disease status. A lod score of 4.08 was obtained with a dominant model where the allele frequency of the susceptibility gene was assumed to be 5% and carriers of the alteration were assumed to have seven-fold the risk of a non-
30 carrier. By inspecting the individual families, no obvious correlation was seen between families that contribute positively to the linkage results with the prevalence of hypertension, diabetes or hyperlipidemias. When the data were reanalyzed with

the hemorrhagic stroke patients removed, the allele sharing lod score increased to 4.86 at D5S2080. Although this 0.46 increase in log score suggests that *STRK1* is involved primarily in ischemic stroke and TIAs, it is not statistically significant based on simulations (one sided P equals 0.09). In order to assess whether such a
5 change in lod score would be likely to occur by chance we selected 1000 random sets of 22 patients whose status we then changed to "unknown" in an analysis. The P value we present is the fraction of the 1000 simulations which produce a lod score increase at the peak locus equal to or greater than that which we observed by changing the affection status of the 22 hemorrhagic stroke patients to "unknown".

10

Identification of Allelic Association

All microsatellite markers in the approx. 6 cM interval (markers from D5S398 to D5S1474) were analyzed with respect to allelic association.

15 *Microsatellite allelic association*

We initially genotyped 864 Icelandic stroke patients and 908 controls using a total of 98 microsatellite markers. These markers are distributed over a region of approximately 11 Mb. The region is centered on our linkage peak and corresponds to the 2 LOD drop. The density of markers is greater in the central 3.7 Mb portion of
20 the region, which includes the 1 LOD drop, with an average spacing of one marker every 53 kb. We have designated this central region, which is flanked by markers D5S1474 and D5S398, as the *STRK1* interval. Three markers, AC027322-5, D5S2121 and AC008818-1, showed a difference in allelic frequency between patients and controls with p-values less than 0.01 (Table 1). Correcting for the
25 relatedness of the Icelandic patients had little impact on the p-values, but after correcting for the number of markers and alleles tested none of these p-values were significant (Table 1).

We had previously observed that our linkage peak increased, albeit not significantly, when excluding the hemorrhagic stroke patients. We therefore tested
30 only those patients with ischemic stroke or TIA for association to the markers. In addition, our ischemic stroke and TIA patients have been sub-classified according to the TOAST research criteria and we also repeated the association analysis separately

for patients with the three TOAST subcategories: cardiogenic, carotid (greater than 70% stenosis) and small vessel occlusive disease. Lastly, we tested the combination of patients with cardiogenic and carotid stroke, since these categories of stroke are most clearly related to atherosclerosis. The results for each of these association studies are presented in Table 1. Three of the tests, one for cardiogenic stroke (AC008818-1), one for carotid stroke (DG5S397), and one for the combination of carotid and cardiogenic stroke (AC008818-1) were significant even after correcting for multiple testing (Table 1). The marker DG5S397 is located within the *PDE4D* gene and AC008818-1 is in the 5' end of *PDE4D* and in the overlapping gene Prostate androgen-regulated transcript (*PART1*) whose transcript is on the other strand going in the opposite direction. *PDE4D* is an important regulator of intracellular levels of cAMP and is expressed widely. *PART1* encodes a putative protein with unknown function predominantly expressed in the prostate gland and in several cancer cell lines. Physical locations of all genotyped markers and *PDE4D* and *PART1* exons are available in Table 2C. The association results for the combination of carotid and cardiogenic stroke were particularly striking with an allele frequency of 35.5% in patients for allele 0 (the CEPH reference allele) of marker AC008818-1 versus 25.5% in controls. The unadjusted p-value for this marker is 0.0000015, and after adjusting for multiple testing of markers is 0.00025 (Table 1). This remains significant even after adjusting for the several phenotypes studied. The risk of this allele to the other alleles of this marker, assuming the multiplicative model Terwilliger, J.D. & Ott, J. A haplotype-based 'haplotype relative risk' approach to detecting allelic associations. *Hum Hered* 42, 337-46 (1992) and Falk, C.T. & Rubinstein, P. Haplotype relative risks: an easy reliable way to construct a proper control sample for risk calculations. *Ann Hum Genet* 51 (Pt 3), 227-33 (1987), was estimated to be 1.60, and the corresponding population attributable risk was 25%.

Thus, the strong association signals from our initial microsatellite association studies helped to focus our attention on the *STRK1* interval and, in particular, to the *PDE4D* gene region.

Table 1.

Microsatellite allelic association analysis of the two-lod drop of the *STRK1* locus.

5 All microsatellites that show association with a p-value less than 0.01 for all stroke, all stroke excluding hemorrhagic stroke, cardiogenic stroke, carotid stroke, small vessel disease and

| Phenotype | Marker | Allele | p-value | RR | # Aff. | Aff. % | # Ctrl | Ctrl. % |
|--|------------|--------|------------|------|--------|-----------|--------|------------|
| All | AC027322-5 | 10 | 0.001 | 3.34 | 787 | 1.90 | 779 | 0.6 |
| | D5S2121 | -2 | 0.0027 | 2.19 | 824 | 2.7 | 870 | 1.3 |
| | AC008818-1 | 0 | 0.0045 | 1.25 | 815 | 29.9 | 891 | 25.5 |
| All patients - excluding hemorrhagic stroke | AC027322-5 | 10 | 0.00052 | 3.56 | 740 | 20 | 779 | 0.6 |
| | D5S2121 | -2 | 0.0023 | 2.23 | 774 | 2.8 | 870 | 1.3 |
| | AC008818-1 | 0 | 0.0062 | 1.24 | 764 | 29.9 | 891 | 25.5 |
| Cardiogenic stroke | AC008818-1 | 0 | 0.000054* | 1.60 | 216 | 35.4 | 891 | 25.5 |
| | D5S1990 | 20 | 0.00053 | 2.18 | 223 | 7.9 | 879 | 3.8 |
| | D5S2089 | -10 | 0.0027 | 2.22 | 219 | 5.9 | 813 | 2.8 |
| | D5S1359 | 2 | 0.0044 | 1.39 | 214 | 36.0 | 777 | 28.8 |
| | AC016604-2 | 0 | 0.0048 | 1.44 | 170 | 51.8 | 446 | 42.7 |
| | AC008804-1 | 0 | 0.0068 | 1.52 | 128 | 36.3 | 367 | 27.3 |
| | AC022125-3 | 0 | 0.0077 | 1.36 | 223 | 36.8 | 775 | 30.0 |
| | DG5S2066 | 0 | 0.0095 | 1.80 | 166 | 92.5 | 501 | 87.2 |
| | DG5S2039 | 9 | 0.0084 | 2.00 | 167 | 8.7 | 491 | 4.6 |
| | D5S647 | -6 | 0.0091 | 2.43 | 199 | 3.8 | 789 | 1.6 |
| Carotid stroke | DG5S397 | 4 | 0.00024* | 1.70 | 124 | 65.7 | 577 | 53.0 |
| | DG5S2056 | 12 | 0.0009 | 3.33 | 80 | 8.8 | 464 | 2.8 |
| | AC008818-1 | 0 | 0.001 | 1.61 | 125 | 35.6 | 891 | 25.5 |
| | DG5S2039 | -3 | 0.003 | 1.62 | 96 | 45.8 | 491 | 34.3 |
| | DG5S2045 | 0 | 0.0051 | 1.80 | 55 | 57.3 | 339 | 42.6 |
| | DG5S818 | 6 | 0.0079 | 1.50 | 111 | 63.1 | 563 | 53.3 |
| | AC016604-3 | 4 | 0.0072 | 1.53 | 99 | 40.9 | 645 | 31.2 |
| Small vessel disease | D5S1359 | 2 | 0.0085 | 1.41 | 157 | 36.3 | 777 | 28.8 |
| | D5S2080 | 2 | 0.0092 | 1.38 | 153 | 54.6 | 885 | 46.5 |
| | D5S2121 | -2 | 0.0059 | 2.93 | 152 | 3.6 | 870 | 1.3 |
| Combined cardiogenic & carotid stroke | AC008818-1 | 0 | 0.0000015* | 1.60 | 341 | 35.5 | 891 | 25.5 |
| | AC008833-6 | 0 | 0.0026 | 1.35 | 335 | 70.3 | 868 | 63.8 |
| | DG5S2066 | 0 | 0.0032 | 1.74 | 258 | 92.3 | 501 | 87.2 |
| | DG5S397 | 4 | 0.009 | 1.29 | 345 | 59.3 | 577 | 53.0 |
| | D5S2121 | -2 | 0.0081 | 2.39 | 336 | 3.0 | 870 | 1.3 |

the combination of cardiogenic and carotid stroke

*significant after adjusting for multiple testing

Alleles #'s: For SNP alleles A = 0, C = 1, G = 2, T = 3; for microsatellite alleles: the CEPH sample 1347-02 (CEPH genomics repository) is used as a reference, the lower allele of each microsatellite in this sample is set at 0 and all other alleles in other samples are numbered accordingly in relation to this reference. Thus allele1 is 1 bp longer than the lower allele in the CEPH sample 1347-02, allele 2 is 2 bp longer than the lower allele in the CEPH sample 1347-02, allele 3 is 3 bp longer than the lower allele in the CEPH sample 1347-02, allele 4 is 4 bp longer than the lower allele in the CEPH sample 1347-02, allele -1 is 1 bp shorter than the lower allele in the CEPH sample 1347-02, allele -2 is 2 bp shorter than the lower allele in the CEPH sample 1347-02, and so on. Note that this same CEPH sample is a standard that is widely used throughout the world for calibration and comparison of alleles.

AC008818-1 amplimer:

TGCTTGGTGAAGGAATAGCCACCCCAGAGAAGGAGTATGGACTTC
TATACACAATCATTTCATTTCATTTCATTTCATTTCATTTCATTTCATTTC
ACTACTCATGCATGATCTTTGTCCTTATCTTCCTCCACTGTCACATGAATA
CCCACCCACTGCACCTACCTGCTTCCTATTCCTGAGAACCCAGGCTC (SEQ
ID NO: 86)

AC008818-1, allele 0 is the same allele as the minimum allele observed in CEPH 1347-02, family 137, individual 02.

Swedish patients have also been genotyped and microsatellite single and multimarker association has been analyzed using the E-M algorithm. A total number of 943 Swedish patients (stroke patients and patients with carotid stenosis) and 322 Swedish controls were analyzed (results shown in Table 2A). At least three haplotypes were more common in patients compared to controls, confirming in a second population that PDE4D shows association to stroke.

Table 2A
Swedish Patient Association

| Markers | Alleles | pAllelic | All Frq Aff | All Frq Ctrl | # aff | # ctrl |
|---|------------|----------|-------------|--------------|-------|--------|
| Swedish patients (n=943) | | | | | | |
| D5S2000 | 2 | 0.0024 | | | 912 | 318 |
| (Sw 2) AC022125-3 AC008833-6 D5S2000 D5S2091 | 0 0 2 0 | 0.006 | 0.035 | 0.01 | 717 | 284 |
| (Sw-1) AC008804-2 D17-H D17-G D5S2080 | -2 4 -2 10 | 0.0028 | 0.057 | 0.05 | 672 | 113 |
| AC008804-2 D17-H D17-G | -4 0 -2 | 0.0037 | 0.056 | 0.03 | 700 | 123 |

Screening for polymorphisms in PDE4D

We next considered whether a functional variant in the *PDE4D* gene might be the cause of our observed microsatellite association. We matched public domain ESTs and our own RT-PCR and RACE transcripts to our sequence of the *STRK1* interval. We defined new alternative *PDE4D* transcripts, which together with previously known transcripts indicated that the *PDE4D* gene contains 22 exons over at least 1.5 Mb and overlaps with *PART1*. The *PDE4D* gene encodes eight protein isoforms and has at least seven promoters. All isoforms identified have an identical C-terminal catalytic domain but differ at the N-terminal regulatory domain (FIG. 2).

We then attempted to identify mutations by sequencing all known *PDE4D* exons (including the overlapping *PART1* exons) and, on average, 100 bp of their flanking introns in 188 patients and 94 controls. Forty-six polymorphisms were identified; 44 SNPs and two intronic deletions. Only two of the polymorphisms, both SNPs, were found within the coding exons of the *PDE4D* gene, which is consistent with the extraordinary lack of variation that others have reported for all four PDE4 classes Houslay, M.D. & Adams, D.R. PDE4 cAMP phosphodiesterases: modular enzymes that orchestrate signalling cross-talk, desensitization and compartmentalization. *Biochem J* 370, 1-18 (2003). The two coding SNPs were typed for additional patients and controls. However, these SNPs did not show significant association to stroke (Table 2B). Therefore, if a functional variant conferring risk for stroke exists in the *PDE4D* gene, it may be within regulatory regions affecting transcription, splicing, message stability, or message transport of one or more isoforms, or in exons that we have not yet identified.

Table 2B

Frequency of *PDE4D* coding mutations.

| Markers | AA change | PDE4D | | p-value | Aff. % | Ctrl. % | # Aff | # Ctrl |
|---------|-----------|-------|--------|---------|--------|---------|-------|--------|
| | | exon | Allele | | | | | |
| SNP 250 | Pro > Thr | D1/D2 | A | 0.163 | 2.0 | 1.5 | 604 | 369 |
| SNP 257 | Lys > Thr | 4 | C | 0.381 | 0.2 | 0.0 | 474 | 294 |

PDE4D isoform expression

Failing to find a functional mutation in the known coding exons of *PDE4D*, we were interested to consider other possible evidence in favor of this gene being a source of the underlying association in this region. We conducted an experiment to study the expression levels of the various isoforms – with any significant differences between patients and controls potentially indicating that regulation of *PDE4D* is a key element in stroke susceptibility. We used EBV transformed B cell lines from randomly selected patients having ischemic stroke or TIA and from controls. We carried out isoform-specific kinetic RT-PCR analysis to quantify each isoform in 83 stroke patients and 84 controls. The patients were principally ischemic stroke patients, with 32 of them having cardiogenic or carotid stroke. We observed that the total *PDE4D* message level, as assessed by amplification across exons present in all isoforms (PAN), was significantly lower in patients than in controls (p-value = 0.0021). This decrease was due primarily to lower expression of the *PDE4D1*, *PDE4D2* and *PDE4D5* isoforms. This significant dysregulation of the expression of multiple *PDE4D* isoforms greatly encouraged us to continue our investigations into the association of the *PDE4D* gene to stroke.

Table 2C: *SNP identification, single marker association and LD mapping of the PDE4D region*

| SNP code | marker or exon | Public name | start in NCBI build 31 | end in NCBI build 33 | start in SEQ ID NO: 1 | end in SEQ ID NO: 1 |
|----------|----------------|-------------|------------------------|----------------------|-----------------------|---------------------|
| | AC016604 - 3 | | 57547045 | 57547304 | | |
| | AC016604 - 2 | | 57623148 | 57623287 | | |
| | exon 11 | | 58241020 | 58241432 | 1655335 | 1655747 |
| | exon 10 | | 58242009 | 58242191 | 1654576 | 1654758 |
| | exon 9 | | 58242702 | 58242824 | 1653943 | 1654065 |
| | exon 8 | | 58243543 | 58243697 | 1653070 | 1653224 |
| | exon 7 | | 58254845 | 58254944 | 1641818 | 1641917 |
| | exon 6 | | 58256107 | 58256271 | 1640491 | 1640655 |
| | exon 5 | | 58257156 | 58257254 | 1639508 | 1639606 |
| | exon 4 | | 58258185 | 58258356 | 1638406 | 1638578 |
| | exon 3 | | 58259724 | 58259817 | 1636944 | 1637037 |
| | exon D1/D2 | | 58305211 | 58305581 | 1591172 | 1591425 |
| | exon LF4 | | 58446946 | 58446995 | 1449835 | 1449884 |
| | exon LF3 | | 58451540 | 58451613 | 1445217 | 1445290 |
| | exon LF2 | | 58459851 | 58459887 | 1436943 | 1436979 |
| | exon LF1 | | 58482128 | 58482319 | 1414511 | 1414702 |
| | AC022125 - 3 | | 58504109 | 58504274 | 1392556 | 1392721 |
| SNP 204 | SNP5PD890407 | | 58506423 | 58506423 | 1390407 | 1390407 |
| | AC008833 - 6 | | 58507019 | 58507222 | 1389608 | 1389811 |
| | exon D9 | | 58541689 | 58542470 | 1354347 | 1355128 |
| | D5S2000 | | 58585460 | 58585849 | | |
| | D5S2091 | | 58593284 | 58593634 | | |
| | exon D8 | | 58623109 | 58623414 | 1273404 | 1273709 |
| | D17 - C | | 58645088 | 58645386 | 1251432 | 1251730 |
| | AC008804 - 1 | | 58784449 | 58784641 | 1112181 | 1112373 |
| | AC008804 - 2 | | 58817743 | 58817931 | 1078881 | 1079069 |
| | exon D3 | | 58852680 | 58852819 | 1044051 | 1044190 |
| | D17 - H | | 58860588 | 58860725 | 1036142 | 1036279 |
| | D17 - G | | 58942270 | 58942541 | 954298 | 954569 |
| | D5S2080 | | 58998685 | 58999021 | | |
| | exon D5 | | 59034598 | 59035009 | 861791 | 862202 |
| | AC027322 - 5 | | 59159221 | 59159326 | 737420 | 737519 |
| | exon D4 | | 59159520 | 59160492 | 736254 | 737226 |
| SNP 102 | SNP5PD166822 | rs714291 | 59229897 | 59229897 | | |
| | exon D7 - 3 | | 59254840 | 59255069 | 641649 | 641878 |
| SNP 101 | SNP5PD138604 | rs1347401 | 59258113 | 59258113 | 638605 | 638605 |
| SNP 100 | SNP5PD121753 | rs1545070 | 59274962 | 59274962 | | |
| SNP 99 | SNP5PD118378 | rs1533019 | 59278338 | 59278338 | | |
| SNP 98 | SNP5PD117029 | rs952110 | 59279687 | 59279687 | | |
| SNP 97 | SNP5PD104361 | rs1995780 | 59292356 | 59292356 | | |
| SNP 96 | SNP5PD97409 | | 59299308 | 59299308 | | |
| SNP 95 | SNP5PD97281 | rs2016324 | 59299437 | 59299437 | | |
| SNP 94 | SNP5PD75406 | rs1396474 | 59321313 | 59321313 | | |
| SNP 93 | SNP5PD73383 | rs1508864 | 59323336 | 59323336 | | |
| SNP 92 | SNP5PD72097 | rs1508859 | 59324622 | 59324622 | | |
| | DG5S2045 | | 59325313 | 59325563 | 571152 | 571406 |
| | DG5S2039 | | 59332799 | 59333077 | 563636 | 563921 |
| SNP 91 | SNP5PD46864 | rs1508863 | 59349851 | 59349851 | | |
| SNP 90 | SNP5PD43868 | rs2136203 | 59352849 | 59352849 | | |
| SNP 89 | SNP5PD29517 | rs1396476 | 59367167 | 59367167 | | |

| | | | | | | |
|--------|---------------|-----------|----------|----------|--------|--------|
| | DG5S2056 | | 59381102 | 59381367 | 515317 | 515582 |
| | DG5S818 | | 59384776 | 59384999 | 511685 | 511908 |
| SNP 88 | SNP5PDM14337 | rs1544788 | 59411021 | 59411021 | | |
| | DG5S397 | | 59438506 | 59438784 | 457900 | 458178 |
| SNP 87 | SNP5PDM43741 | rs2910829 | 59440424 | 59440424 | | |
| | exon D7 - 2 | | 59451909 | 59452039 | 444645 | 444775 |
| SNP 86 | SNP5PDM57997 | rs2962972 | 59454680 | 59454680 | | |
| SNP 85 | SNP5PDM65461 | rs2961897 | 59462144 | 59462144 | | |
| SNP 84 | SNP5PDM67604 | rs719702 | 59464287 | 59464287 | | |
| SNP 83 | SNP5PDM76361 | rs966221 | 59473045 | 59473045 | | |
| SNP 82 | SNP5PDM83539 | rs2961903 | 59480223 | 59480223 | | |
| SNP 81 | SNP5PDM89176 | | 59485859 | 59485859 | 410826 | 410826 |
| SNP 80 | SNP5PDM89683 | rs1862614 | 59486368 | 59486368 | | |
| | DG5S2066 | | 59522085 | 59522346 | 374339 | 374600 |
| SNP 79 | SNP5PDM132154 | | 59528838 | 59528838 | 367847 | 367847 |
| SNP 78 | SNP5PDM153120 | | 59549804 | 59549804 | 346881 | 346881 |
| SNP 77 | SNP5PDM161561 | | 59558245 | 59558245 | 338440 | 338440 |
| SNP 76 | SNP5PDM166786 | | 59563470 | 59563470 | 333215 | 333215 |
| SNP 75 | SNP5PDM181173 | | 59577856 | 59577856 | 318829 | 318829 |
| SNP 74 | SNP5PDM182792 | | 59579475 | 59579475 | 317210 | 317210 |
| SNP 73 | SNP5PDM211974 | | 59608650 | 59608650 | 288027 | 288027 |
| SNP 72 | SNP5PDM217886 | | 59614557 | 59614557 | 282115 | 282115 |
| SNP 71 | SNP5PDM218639 | | 59615310 | 59615310 | 281362 | 281362 |
| SNP 70 | SNP5PDM224528 | | 59621190 | 59621190 | 275473 | 275473 |
| SNP 69 | SNP5PDM236461 | rs1423248 | 59633124 | 59633124 | | |
| SNP 68 | SNP5PDM259844 | | 59656504 | 59656504 | 240157 | 240157 |
| SNP 67 | SNP5PDM261488 | | 59658148 | 59658148 | 238513 | 238513 |
| SNP 66 | SNP5PDM265669 | | 59662328 | 59662328 | 234332 | 234332 |
| SNP 65 | SNP5PDM271674 | rs918590 | 59668333 | 59668333 | | |
| SNP 64 | SNP5PDM275805 | rs1423247 | 59672463 | 59672463 | | |
| SNP 63 | SNP5PDM280894 | rs789389 | 59677551 | 59677551 | | |
| SNP 62 | SNP5PDM285592 | | 59682247 | 59682247 | 214409 | 214409 |
| SNP 61 | SNP5PDM296955 | rs37691 | 59693610 | 59693610 | | |
| SNP 60 | SNP5PDM299842 | | 59696497 | 59696497 | 200159 | 200159 |
| SNP 59 | SNP5PDM307243 | rs37684 | 59703890 | 59703890 | | |
| SNP 58 | SNP5PDM308509 | rs2898278 | 59705155 | 59705155 | | |
| SNP 57 | SNP5PDM310220 | rs401207 | 59706866 | 59706866 | | |
| SNP 56 | SNP5PDM310653 | rs702553 | 59707298 | 59707298 | | |
| SNP 55 | SNP5PDM324741 | rs251726 | 59721387 | 59721387 | | |
| SNP 54 | SNP5PDM326519 | rs27223 | 59723165 | 59723165 | | |
| SNP 53 | SNP5PDM329913 | | 59726556 | 59726556 | 170088 | 170088 |
| SNP 52 | SNP5PDM332989 | | 59729632 | 59729632 | 166900 | 166900 |
| SNP 51 | SNP5PDM338487 | | 59735122 | 59735122 | 161514 | 161514 |
| SNP 50 | SNP5PDM345627 | rs173591 | 59742248 | 59742248 | | |
| SNP 49 | SNP5PDM349039 | rs27220 | 59745661 | 59745661 | | |
| SNP 48 | SNP5PDM351840 | rs37760 | 59748461 | 59748461 | | |
| SNP 47 | SNP5PDM356081 | | 59752701 | 59752701 | 143922 | 143922 |
| SNP 46 | SNP5PDM356447 | | 59753067 | 59753067 | 143555 | 143555 |
| SNP 45 | SNP5PDM357221 | | 59753842 | 59753842 | 142780 | 142780 |
| SNP 44 | SNP5PDM357245 | | 59753865 | 59753865 | 142757 | 142757 |
| SNP 43 | SNP5PDM357445 | | 59754066 | 59754066 | 142556 | 142556 |
| | PART1-exon 1 | | 59754284 | 59754775 | | |
| | exon D7 - 1 | | 59754294 | 59754415 | 142207 | 142328 |

| | | | | | | |
|--------|---------------|------------------|----------|----------|--------|--------|
| | PART1-exon 2 | | 59756013 | 59757617 | | |
| SNP 42 | SNP5PDM361194 | rs153031 | 59757816 | 59757816 | | |
| SNP 41 | SNP5PDM361545 | | 59758341 | 59758341 | 138456 | 138456 |
| | AC008818 - 1 | SEQ ID NO: 86 | 59759882 | 59760075 | 136740 | 136547 |
| SNP 40 | SNP5PDM363736 | | 59760357 | 59760357 | 136265 | 136265 |
| SNP 39 | SNP5PDM364360 | rs3887175 | 59760981 | 59760981 | | |
| SNP 38 | SNP5PDM364848 | | 59761469 | 59761469 | 135152 | 135152 |
| SNP 37 | SNP5PDM364888 | rs26956 | 59761510 | 59761510 | 135112 | 135112 |
| SNP 36 | SNP5PDM366629 | | 59763250 | 59763250 | 133371 | 133371 |
| SNP 35 | SNP5PDM367438 | rs26955 | 59764060 | 59764060 | 132562 | 132562 |
| SNP 34 | SNP5PDM368135 | rs27653 | 59764755 | 59764755 | 131865 | 131865 |
| SNP 33 | SNP5PDM369610 | | 59766229 | 59766229 | 130391 | 130391 |
| SNP 32 | SNP5PDM370640 | | 59767259 | 59767259 | 129361 | 129361 |
| SNP 31 | SNP5PDM370641 | rs457053 | 59767260 | 59767261 | 129360 | 129360 |
| SNP 30 | SNP5PDM374696 | rs27221 | 59771316 | 59771316 | 125304 | 125304 |
| SNP 29 | SNP5PDM376181 | rs2963110 | 59772800 | 59772800 | | |
| SNP 28 | SNP5PDM376575 | rs35387 | 59773194 | 59773194 | 123426 | 123426 |
| SNP 27 | SNP5PDM376688 | rs35386 | 59773308 | 59773308 | 123312 | 123312 |
| SNP 26 | SNP5PDM379372 | rs40512 | 59775992 | 59775992 | 120628 | 120628 |
| SNP 25 | SNP5PDM380376 | | 59776995 | 59776995 | | |
| SNP 24 | SNP5PDM381086 | rs35385 | 59777706 | 59777706 | 118914 | 118914 |
| SNP 23 | SNP5PDM388220 | rs26953 | 59784839 | 59784839 | 111781 | 111781 |
| SNP 22 | SNP5PDM388748 | | 59785368 | 59785368 | 111252 | 111252 |
| SNP 21 | SNP5PDM388749 | rs26954 | 59785369 | 59785370 | | |
| SNP 20 | SNP5PDM390700 | | 59787319 | 59787319 | 109301 | 109301 |
| SNP 19 | SNP5PDM392152 | rs4133470 | 59788771 | 59788771 | 107849 | 107849 |
| SNP 18 | SNP5PDM392684 | | 59789302 | 59789302 | 107317 | 107317 |
| SNP 17 | SNP5PDM394085 | | 59790704 | 59790704 | 105792 | 105792 |
| SNP 16 | SNP5PDM394776 | rs35384 | 59791395 | 59791395 | 105225 | 105225 |
| SNP 15 | SNP5PDM395449 | rs35382 | 59792068 | 59792068 | 104552 | 104552 |
| SNP 14 | SNP5PDM397023 | rs26950 | 59793643 | 59793643 | 102977 | 102977 |
| SNP 13 | SNP5PDM399206 | rs26949 | 59795825 | 59795825 | 100795 | 100795 |
| SNP 12 | SNP5PDM400966 | rs153153 | 59797585 | 59797585 | 99035 | 99035 |
| SNP 11 | SNP5PDM402736 | rs152340 | 59799349 | 59799349 | | |
| SNP 10 | SNP5PDM407853 | | 59804468 | 59804468 | 92148 | 92148 |
| SNP 9 | SNP5PDM408531 | | 59805145 | 59805145 | 91470 | 91470 |
| SNP 8 | SNP5PDM408979 | | 59805593 | 59805593 | 91022 | 91022 |
| SNP 7 | SNP5PDM409460 | | 59806074 | 59806074 | 90541 | 90541 |
| SNP 6 | SNP5PDM411387 | | 59808001 | 59808001 | 88614 | 88614 |
| SNP 5 | SNP5PDM411544 | rs27564 | 59808159 | 59808159 | 88456 | 88456 |
| SNP 4 | SNP5PDM416882 | rs153152 | 59813496 | 59813496 | 83119 | 83119 |
| SNP 3 | SNP5PDM417756 | rs187481 | 59814371 | 59814371 | 82244 | 82244 |
| SNP 2 | SNP5PDM419874 | rs152341 | 59816488 | 59816488 | 80127 | 80127 |
| SNP 1 | SNP5PDM421449 | rs248911 | 59818063 | 59818063 | 78552 | 78552 |
| | D5S1990 | | 60945599 | 60945816 | | |
| | D5S1359 | | 63542603 | 63542894 | | |
| | D5S2089 | | 65914315 | 65914496 | | |
| | D5S647 | | 66217674 | 66218065 | | |
| | D5S2121 | | 66584091 | 66584385 | | |

We next searched for SNPs in the intronic and flanking regions of *PDE4D*. The SNPs were identified in the public NCBI SNP database or by sequencing selected intronic and flanking regions in the gene in at least 94 patients and 94 controls. We initially identified 637 SNPs. Many of these SNPs were completely correlated so we removed many redundant SNPs from further genotyping. Some SNPs with very low minor allele frequencies were also ignored. This resulted in a set of 260 SNPs that were then genotyped for the entire patient and control cohorts. The preponderance of markers with significant associations was located at the 5' end of the gene. One SNP (SNP5PDM76361;SNP83) for carotid stroke and five of the SNPs (SNP5PDM357221=SNP45, SNP5PDM361545=SNP41, SNP5PDM43741=SNP87, SNP5PDM29517=SNP89 and SNP5PDM56) for the combined cardiogenic and carotid stroke remained significant even after adjusting for all the SNPs tested (Table 2D). Three of these significant SNPs flank exon D7-1; the other three are in a 100 kb region containing exon D7-2 (for physical positions see Table 2D). The two most significant SNPs, SNP45 and SNP41, are within 6 kb of the microsatellite marker AC008818-1, and the at-risk alleles of all three genetic markers are in strong linkage disequilibrium with $D' > 0.9$ and p-value nearly zero. The square of the correlation (R^2) is very high between the two SNPs (~ 0.93), but is substantially lower (~ 0.08) between each SNP and the at-risk allele of the microsatellite. This is due to the fact that the frequency of the at-risk alleles of the two SNPs are similar, and much more frequent than that for the at-risk allele of the microsatellite. The LD block structure around the 5' end of *PDE4D* is displayed in FIG. 13.1. We delineate three blocks A, B and C encompassing the first three exons of *PDE4D* and its immediate downstream region. Exons D7-3 and D7-2 are both in block A, while D7-1 (the first exon) is in block B, but close to its border with block C. Given this block structure we were prepared to investigate the haplotype associated susceptibility to stroke in this region.

Table 2D. All SNPs that show association with a p-value less than 0.01 for all stroke patients, all patients excluding hemorrhagic stroke and the combined cardiogenic and carotid stroke.

| Phenotype | Marker | Allele | p-value | RR | # Affect | Aff. % | # Ctrl | Ctrl. % |
|--|---------|--------|-----------|------|----------|--------|--------|---------|
| All patients | | | | | | | | |
| | SNP 32 | C | 0.00024 | 1.46 | 400 | 37.9 | 475 | 29.5 |
| | SNP 56 | T | 0.0028 | 1.31 | 550 | 71.4 | 615 | 65.5 |
| | SNP 45 | G | 0.0065 | 1.33 | 723 | 82.4 | 492 | 78.0 |
| | SNP 48 | T | 0.0091 | 1.28 | 547 | 68.3 | 481 | 62.8 |
| All patients excl. hemorrhagic stroke | | | | | | | | |
| | SNP 32 | C | 0.00034 | 1.45 | 377 | 37.8 | 475 | 29.5 |
| | SNP 56 | T | 0.0066 | 1.28 | 518 | 70.9 | 615 | 65.5 |
| | SNP 45 | G | 0.0095 | 1.31 | 679 | 82.3 | 492 | 78.0 |
| Combined cardiogenic & carotid | | | | | | | | |
| | SNP 45 | G | 0.000034* | 1.77 | 309 | 86.3 | 492 | 78.0 |
| | SNP 41 | A | 0.000078* | 1.86 | 236 | 86.0 | 368 | 76.8 |
| | SNP 87 | T | 0.00019* | 1.49 | 263 | 58.2 | 583 | 48.4 |
| | SNP 89 | A | 0.00025* | 1.84 | 232 | 88.8 | 450 | 81.1 |
| | SNP 56 | T | 0.00027* | 1.56 | 230 | 74.8 | 615 | 65.5 |
| | SNP 39 | T | 0.00032 | 1.58 | 326 | 84.4 | 589 | 77.3 |
| | SNP 91 | G | 0.00047 | 1.80 | 233 | 88.6 | 451 | 81.3 |
| | SNP 32 | C | 0.00069 | 1.61 | 144 | 40.3 | 475 | 29.5 |
| | SNP 62 | A | 0.00089 | 1.73 | 153 | 83.0 | 556 | 73.8 |
| | SNP 48 | T | 0.00080 | 1.51 | 229 | 71.8 | 481 | 62.8 |
| | SNP 42 | A | 0.0018 | 1.49 | 259 | 72.0 | 403 | 63.6 |
| | SNP 184 | G | 0.0025 | 1.68 | 252 | 90.7 | 570 | 85.3 |
| | SNP 58 | T | 0.0042 | 1.54 | 234 | 85.3 | 569 | 79.0 |
| | SNP 53 | C | 0.0041 | 1.58 | 146 | 36.0 | 269 | 26.2 |
| | SNP 97 | G | 0.0046 | 1.40 | 225 | 54.2 | 450 | 45.9 |
| | SNP 204 | A | 0.0049 | 1.32 | 334 | 63.9 | 651 | 57.3 |
| | SNP 8 | A | 0.0054 | 1.59 | 228 | 89.0 | 612 | 83.7 |
| | SNP 83 | C | 0.0074 | 1.39 | 223 | 60.1 | 349 | 52.0 |
| | SNP 43 | T | 0.0093 | 1.48 | 243 | 85.4 | 550 | 79.8 |

* significant after adjusting for multiple testing

5

Haplotype analysis

Our general approach to haplotype analysis involves using likelihood-based inference applied to NEsted MOdels. The method is implemented in our program NEMO, which allows for many polymorphic markers, SNPs and microsatellites. The

method and software are specifically designed for case-control studies where the purpose is to identify haplotype groups that confer different risks. It is also a tool for studying LD structures.

When investigating haplotypes constructed from many markers, apart from
 5 looking at each haplotype individually, meaningful summaries often require putting haplotypes into groups. A particular partition of the haplotype space is a model that assumes haplotypes within a group have the same risk, while haplotypes in different groups can have different risks. Two models/partitions are nested when one, the alternative model, is a finer partition compared to the other, the null model, *i.e.*, the
 10 alternative model allows some haplotypes assumed to have the same risk in the null model to have different risks. The models are nested in the classical sense that the null model is a special case of the alternative model. Hence traditional generalized likelihood ratio tests can be used to test the null model against the alternative model. Note that, with a multiplicative model, if haplotypes h_i and h_j are assumed to have
 15 the same risk, it corresponds to assuming that $f_i/p_i = f_j/p_j$ where f and p denote haplotype frequencies in the affected population and the control population respectively.

One common way to handle uncertainty in phase and missing genotypes is a two-step method of first estimating haplotype counts and then treating the estimated
 20 counts as the exact counts, a method that can sometimes be problematic (*e.g.*, see the information measure section below) and may require randomization to properly evaluate statistical significance. In NEMO, maximum likelihood estimates, likelihood ratios and p-values are calculated directly, with the aid of the EM algorithm, for the observed data treating it as a missing-data problem.

25 NEMO allows complete flexibility for partitions. For example, the first haplotype problem described in the Methods section on Statistical analysis considers testing whether h_1 has the same risk as the other haplotypes h_2, \dots, h_k . Here the alternative grouping is $[h_1], [h_2, \dots, h_k]$ and the null grouping is $[h_1, \dots, h_k]$. The second haplotype problem in the same section involves three haplotypes $h_1 = G0$, $h_2 = GX$ and $h_3 = AX$, and the focus is on comparing h_1 and h_2 . The alternative
 30 grouping is $[h_1], [h_2], [h_3]$ and the null grouping is $[h_1, h_2], [h_3]$. The actual problem we faced in FIG. 11.1 is actually slightly more complicated because allele X is a

composite allele that includes five alleles other than allele 0, and hence GX and AX each correspond to five haplotypes. One could have collapsed these alleles into one at the data processing stage, and performed the test as described. This is a perfectly valid approach, and indeed, whether we collapse or not makes no difference if there were no missing information regarding phase. But, with the actual data, each of the 5 alleles making up X correlates differently with the SNP alleles and this provides some partial information on phase. Collapsing at the data processing stage will unnecessarily increase the amount of missing information. What was actually done is natural in the nested-models/partition framework. Let h_2 be split into $h_{2a}, h_{2b}, \dots, h_{2e}$, and h_3 be split into $h_{3a}, h_{3b}, \dots, h_{3e}$. Then the alternative grouping is $[h_1], [h_{2a}, h_{2b}, \dots, h_{2e}], [h_{3a}, h_{3b}, \dots, h_{3e}]$ and the null grouping is $[h_1, h_{2a}, h_{2b}, \dots, h_{2e}], [h_{3a}, h_{3b}, \dots, h_{3e}]$. The same method is used to handle the composite haplotypes in FIG. 11.2 and 11.3 where collapsing at the data processing stage is not even an option since L_C represents multiple haplotypes constructed from 25 SNPs. Here, we also want to mention that, apart from the pair-wise comparisons presented in FIG. 11.1, a 3-way test with the alternative grouping of $[h_1], [h_{2a}, h_{2b}, \dots, h_{2e}], [h_{3a}, h_{3b}, \dots, h_{3e}]$ versus the null grouping of $[h_1, h_{2a}, h_{2b}, \dots, h_{2e}, h_{3a}, h_{3b}, \dots, h_{3e}]$ could also be performed. Note that the generalized likelihood ratio test-statistic would have two degrees of freedom instead of one. We actually have performed this test and it gave a p-value of 2.4×10^{-7} .

Measuring information

Even though likelihood ratio tests based on likelihoods computed directly for the observed data, which have captured the information loss due to uncertainty in phase and missing genotypes, can be relied on to give valid p-values, it would still be of interest to know how much information had been lost due to the information being incomplete. Interestingly, one can measure information loss by considering a two-step procedure to evaluating statistical significance that appears natural but happens to be systematically anti-conservative. Suppose we calculate the maximum likelihood estimates for the population haplotype frequencies calculated under the alternative hypothesis that there are differences between the affected population and control population, and use these frequency estimates as estimates of the observed

frequencies of haplotype counts in the affected sample and in the control sample. Suppose we then perform a likelihood ratio test treating these estimated haplotype counts as though they are the actual counts. We could also perform a Fisher's exact test, but we would then need to round off these estimated counts since they are in general non-integers. This test will in general be anti-conservative because treating the estimated counts as if they were exact counts ignores the uncertainty with the counts, overestimates the effective sample size and underestimates the sampling variation. It means that the chi-square likelihood-ratio test statistic calculated this way, denoted by Λ^* , will in general be bigger than Λ , the likelihood-ratio test-statistic calculated directly from the observed data as described in methods. But Λ^* is useful because the ratio Λ/Λ^* happens to be a good measure of information, or $1 - (\Lambda/\Lambda^*)$ is a measure of the fraction of information lost due to missing information. This information measure for haplotype analysis is described in Nicolae and Kong, Technical Report 537, Department of Statistics, University of Statistics, University of Chicago, Revised for *Biometrics* (2003) as a natural extension of information measures defined for linkage analysis, and is implemented in NEMO.

Haplotype association

We first considered haplotypes based on the most significantly associated SNPs and microsatellite, SNP45, SNP41 and AC008818-1, which are all in block B and are separated by only 6 kb. Not surprisingly given the high degree of correlation between SNP45 and SNP41, we found that it was sufficient to consider only the two marker haplotypes consisting of the microsatellite and SNP45 – the SNP with the higher genotype yield. The results of this association study for the combination of carotid and cardiogenic stroke are displayed in FIG. 11.1. Note that, for convenience, we have designated by the letter X the joint set of alleles that are not the at-risk allele, 0, of microsatellite AC008818-1. Thus, GX should be understood as the composite of all haplotypes including the G nucleotide of SNP45 except for the G0 haplotype. For our samples, the A0 haplotype does not exist. This suggests that allele 0 originated in a haplotype background with allele G of SNP45, and since then no recombination has occurred between those two markers for chromosomes that carried allele 0. AX, G0 and GX have significantly distinct risks for the

combined carotid and cardiogenic stroke phenotype. We refer to GX as the wild type because it is the most common (53.4% in controls) and also because it has the intermediate level risk that is not too different from the population risk. The haplotype G0 has increased risk and AX is protective, with risks of 1.46 and 0.70
5 relative to the wild type, respectively. The G0 risk is 2.07 times that of the protective haplotype AX. Each of the three pairwise comparisons is highly significant, with p-values ranging from 0.006 to 7.2×10^{-8} . It is interesting to observe that even though both AX and GX are composite haplotypes, the AX haplotype can be simply summarized by the allele A of SNP45, since the A0 haplotype does not exist. For a
10 similar reason, the G0 haplotype is completely determined by the 0 allele of AC008818-1. Also displayed in FIG. 11.1 is the information content (Info) of each test. The difference between Info and 1 is a measure of the information that is lost due to the uncertainty with phase and missing genotypes. Note that Info is very close to 1 for each of the three tests in FIG. 11.1. That is a result of SNP45 and
15 AC008818-1 being in very strong LD. Note that tests presented later in FIG. 11.2 and 11.3, involving longer haplotypes have lower information content.

We next identified and estimated the risks for the common SNP haplotypes within each block. For this portion of the analysis only those SNPs with minor allele frequency greater than 20% were considered. Block A (300 kb) contained 19 such
20 SNPs, block B (200 kb) 22 SNPs, and block C (60 kb) 25 SNPs. All haplotypes within each block with an estimated frequency in the population of 2% or greater have been identified. Within each block there were fewer than ten such haplotypes, and they accounted for approximately 80% of the total haplotype frequency for that block. A brief schematic of the identified haplotypes are displayed in FIG. 13.2
25 and the risks and frequencies of these haplotypes are available in Table 3. Within block A no common haplotype has greater risk than SNP87 alone. The strongest signals were for haplotypes in block B and C. Each block contained a haplotype significantly associated with the combination of carotid and cardiogenic stroke and having relative risk around 1.5. The common at-risk haplotype in block B is the SNP
30 background of the G0 haplotype previously identified.

While there were no significant single marker associations in block C, a common haplotype with 15.4% frequency in controls was observed. We designate

this haplotype H_C . Investigation of the contribution of H_C in conjunction with the SNP45 and AC008818-1 haplotypes leads to another interesting observation. For notation, all haplotypes defined by the 25 SNPs in block C that are not H_C are jointly denoted by the composite haplotype L_C . First, it is noted that AX and H_C do not exist
5 together on the same chromosome (see FIG. 11.3), at least in these samples, and thus blocks B and C are far from being independent. As a consequence, the extended composite haplotype AXL_C is the same as AX. The haplotype G0 can be split into the two extended haplotypes $G0H_C$ and the composite $G0L_C$, which, as indicated in FIG. 11.2, have significantly different risks (p value = 0.0067). Moreover, it appears
10 that the elevated risk of G0 is totally accounted for by $G0H_C$ as $G0L_C$ has risk that is not significantly different from $GX = GXH_C + GXL_C$ (see FIG. 11.2). This observation allows us to refine the haplotype groupings of FIG. 13.1 into the groupings indicated by FIG. 13.3. The extended at-risk haplotype $G0H_C$ (8.8% in controls) and protective composite haplotype AXL_C (21.1% in controls), have,
15 respectively, relative risks of 1.98 and 0.68 compared to the wild type (70.1% in controls). Based on these risk estimates, if everybody's risk can be made to correspond to that of a homozygote carrier of the protective variant, the number of cases would be reduced by 55%, which can be interpreted as the population attributed risk of the at-risk haplotype and the wild type combined.

20 The at-risk haplotype $G0H_C$ spans a region of about 64kb. While it is possible that the increased risk is due to multiple polymorphisms over that region, the results are also consistent with a relatively recent mutation, as yet to be identified, which occurred in that haplotype background, and since then no recombination has occurred in that extended region for chromosomes carrying the
25 mutation. By contrast, the protective composite haplotype AXL_C can be simply represented by allele A of SNP45. Hence, it is possible that allele A of SNP45 is the functional protective variant, although it is possible that the functional variant is simply in strong LD with allele A of SNP45 and has yet to be identified. Indeed, statistically, the effects of SNP45 and SNP41 are indistinguishable from each other.

Statistical analysis.

For single marker association to the disease, the Fisher exact test was used to calculate two-sided p-values for each individual allele. All p-values were presented unadjusted for multiple comparisons unless specifically indicated. The presented
5 frequencies (for microsatellites, SNPs and haplotypes) were allelic frequencies as opposed to carrier frequencies. To minimize any bias due the relatedness of the patients who were recruited as families for the linkage analysis, we eliminated first and second-degree relatives from the patient list. Furthermore, we have repeated the test for association correcting for any remaining relatedness among the patients, by
10 extending a variance adjustment procedure described in Risch, N. & Teng, J. (*Genome Res.*, 8:1278-1288 (1998)). The relative power of family-based and case-control designs for linkage disequilibrium studies of complex human diseases I. DNA pooling. (*ibid*) for sibships so that it can be applied to general familial relationships, and present both adjusted and unadjusted p-values for comparison. The
15 differences are in general very small as expected. To assess the significance of single-marker association corrected for multiple testing we carried out a randomisation test using the same genotype data. We randomised the cohorts of patients and controls and redid the association analysis. This procedure was repeated up to 500,000 times and the p-value we presented is the fraction of replications that
20 produced a p-value for some marker allele that is lower than or equal to the p-value we observed using the original patient and control cohorts.

For both single-marker and haplotype analyses, relative risk (RR) and the population attributable risk (PAR) were calculated assuming a multiplicative model (haplotype relative risk model), (Terwilliger, J.D. & Ott, J., *Hum Hered*, 42, 337-46
25 (1992) and Falk, C.T. & Rubinstein, P, *Ann Hum Genet* 51 (Pt 3), 227-33 (1987)), i.e., that the risks of the two alleles/haplotypes a person carries multiply. For example, if RR is the risk of A relative to a, then the risk of a person homozygote AA will be RR times that of a heterozygote Aa and RR^2 times that of a homozygote aa. The multiplicative model has a nice property that simplifies analysis and
30 computations — haplotypes are independent, i.e., in Hardy-Weinberg equilibrium, within the affected population as well as within the control population. As a consequence, haplotype counts of the affecteds and controls each have multinomial

distributions, but with different haplotype frequencies under the alternative hypothesis. Specifically, for two haplotypes h_i and h_j , $\text{risk}(h_i)/\text{risk}(h_j) = (f_i/p_i)/(f_j/p_j)$, where f and p denote respectively frequencies in the affected population and in the control population. While there is some power loss if the true model is not
 5 multiplicative, the loss tends to be mild except for extreme cases. Most importantly, p-values are always valid since they are computed with respect to null hypothesis.

In general, haplotype frequencies are estimated by maximum likelihood and tests of differences between cases and controls are performed using a generalized likelihood ratio test (Rice, J.A. *Mathematical Statistics and Data Analysis*, 602
 10 (International Thomson Publishing, (1995)). deCODE's haplotype analysis program called NEMO, which stands for NEsted MOdels, was used to calculate all the haplotype results presented. To handle uncertainties with phase and missing genotypes, it is emphasized that we do not use a common two-step approach to association tests, where haplotype counts are first estimated, possibly with the use of
 15 the EM algorithm, Dempster, (A.P., Laird, N.M. & Rubin, D.B., *Journal of the Royal Statistical Society B*, 39, 1-38 (1971)) and then tests are performed treating the estimated counts as though they are true counts, a method that can sometimes be problematic and may require randomisation to properly evaluate statistical significance. Instead, with NEMO, maximum likelihood estimates, likelihood ratios
 20 and p-values are computed with the aid of the EM-algorithm directly for the observed data, and hence the loss of information due to uncertainty with phase and missing genotypes is automatically captured by the likelihood ratios. Even so, it is of interest to know how much information is retained, or lost, due to incomplete information. Described herein is such a measure that is natural under the likelihood
 25 framework. For a fixed set of markers, the simplest tests we performed, with results presented in Table 3, compare one selected haplotype against all the others. Call the selected haplotype h_1 and the others h_2, \dots, h_k . Let p_1, \dots, p_k denote the population frequencies of the haplotypes in the controls, and f_1, \dots, f_k denote the population frequencies of the haplotypes in the affecteds. Under the null hypothesis, $f_i = p_i$ for
 30 all i . The alternative model we use for the test assumes h_2, \dots, h_k to have the same risk while h_1 is allowed to have a different risk. This implies that while p_1 can be different from f_1 , $f_i/(f_2 + \dots + f_k) = p_i/(p_2 + \dots + p_k) = \beta_i$ for $i = 2, \dots, k$. Denoting f_1/p_1 by

r , and noting that $\beta_2 + \dots + \beta_k = 1$, the test statistic based on generalized likelihood ratios is

$$\Lambda = 2 \left[\ell(\hat{r}, \hat{p}_1, \hat{\beta}_2, \dots, \hat{\beta}_{k-1}) - \ell(1, \tilde{p}_1, \tilde{\beta}_2, \dots, \tilde{\beta}_{k-1}) \right]$$

where ℓ denotes log_elikelihood and \sim and \wedge denote maximum likelihood estimates under the null hypothesis and alternative hypothesis respectively. Λ has asymptotically a chi-square distribution with 1-df, under the null hypothesis and it was used to compute p-values presented in Table 3. The tests presented in FIG. 11 have slightly more complicated null and alternative hypotheses. For the results in FIG. 11, let h_1 be G0, h_2 be GX and h_3 be AX. When comparing G0 against GX, *i.e.*, this is the test which gives estimated RR of 1.46 and p-value = 0.0002, the null assumes G0 and GX have the same risk but AX is allowed to have a different risk. The alternative hypothesis allows all three haplotype groups to have different risks. This implies that, under the null hypothesis, there is a constraint that $f_1/p_1 = f_2/p_2$, or $w = [f_1/p_1]/[f_2/p_2] = 1$. The test statistic based on generalized likelihood ratios is

$$\Lambda = 2 \left[\ell(\hat{p}_1, \hat{f}_1, \hat{p}_2, \hat{w}) - \ell(\tilde{p}_1, \tilde{f}_1, \tilde{p}_2, 1) \right]$$

that again has asymptotically a chi-square distribution with 1-df under the null hypothesis. There is actually an extra complication to the test due to h_2 and h_3 being composite haplotypes. That is handled in a natural manner under the nested models framework. Other tests presented in FIG. 11.2 and 11.3 were similarly performed.

LD between pairs of SNPs was calculated using the standard definition of D' and R^2 (Lewontin, R., *Genetics* 49, 49-67 (1964) and Hill, W.G. & Robertson, A. *Theor. Appl. Genet.* 22, 226-231 (1968)). Using NEMO, frequencies of the two marker allele combinations are estimated by maximum likelihood and deviation from linkage equilibrium is evaluated by a likelihood ratio test. The definitions of D' and R^2 were extended to include microsatellites by averaging over the values for all possible allele combination of the two markers weighted by the marginal allele probabilities. When plotting all marker combination to elucidate the LD structure in a particular region, we plot D' in the upper left corner and the p-value in the lower right corner. In the LD plots we present the markers are plotted equidistant rather than according to their physical location.

Table 3 Haplotype diversity at the 5' end of the PDE4D gene.

All haplotypes shown that have > 2% population frequency within each of the 3 blocks of strong LD together with the haplotype association results comparing the combination of cardiogenic and carotid stroke versus controls.

5

Block A:

| SNP 102 | SNP 101 | SNP 100 | SNP 99 | SNP 98 | SNP 97 | SNP 96 | SNP 95 | SNP 93 | SNP 92 | SNP 90 | SNP 88 | SNP 87 | SNP 86 | SNP 84 | SNP 83 | SNP 82 | SNP 81 | SNP 80 | SNP 79 | p-value | Aff % | Ctrl % | RR |
|---------|---------|---------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|---------|-------|--------|------|
| T | G | T | A | T | G | A | G | A | G | A | G | T | A | G | C | G | G | T | C | 0.0302 | 26.7 | 22.2 | 1.28 |
| C | G | C | A | C | C | G | A | G | A | G | A | C | G | G | C | G | G | T | C | 0.366 | 2.1 | 2.9 | 0.73 |
| C | G | C | A | C | C | G | A | G | A | G | A | C | G | A | T | G | G | A | T | 0.303 | 2.6 | 3.6 | 0.71 |
| C | G | C | A | C | C | G | A | G | A | G | A | C | G | A | T | A | G | A | T | 0.335 | 22.5 | 24.5 | 0.89 |
| C | G | C | G | C | G | A | G | A | G | A | G | T | A | G | C | G | A | A | T | 0.216 | 12.6 | 10.6 | 1.23 |
| C | A | C | A | C | C | A | G | A | G | A | G | T | A | G | C | G | A | A | T | 0.876 | 2.2 | 2.4 | 0.94 |
| C | A | C | A | C | C | A | G | A | G | A | G | C | A | A | T | G | G | T | C | 0.001 | 6.5 | 11.2 | 0.55 |
| C | A | C | G | C | G | A | G | A | G | A | G | T | A | G | C | G | A | A | T | 0.401 | 7.9 | 6.6 | 1.20 |

10 Block B:

| SNP-77 | SNP-76 | SNP-73 | SNP-71 | SNP-69 | SNP-67 | SNP-66 | SNP-64 | SNP-63 | SNP-62 | SNP-61 | SNP-59 | SNP-57 | SNP-56 | SNP-54 | SNP-53 | SNP-49 | SNP-48 | SNP-45 | SNP-42 | SNP-41 | SNP-39 | p-value | Aff % | Ctrl % | RR |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|---------|-------|--------|------|
| A | A | T | G | T | A | A | G | A | A | C | A | G | T | A | C | C | T | G | A | A | T | 0.0004 | 29.2 | 21.4 | 1.52 |
| A | A | T | G | T | A | A | G | A | C | T | A | A | A | A | T | T | C | A | G | G | A | 0.007 | 4.3 | 7.6 | 0.55 |
| G | A | C | A | T | A | A | G | A | A | C | A | G | T | A | C | C | T | G | A | A | T | 0.958 | 2.0 | 2.1 | 0.98 |
| G | A | C | A | T | G | G | A | G | A | T | A | A | A | A | T | T | C | G | G | A | T | 0.610 | 6.2 | 5.6 | 1.13 |
| G | A | C | A | T | G | G | A | G | C | T | A | A | A | A | T | T | C | A | G | G | A | 0.0104 | 3.4 | 6.3 | 0.52 |
| G | G | C | A | T | G | A | G | A | A | C | C | G | T | G | T | C | T | G | A | A | T | 0.803 | 14.6 | 14.1 | 1.04 |

Block C:

15

| SNP 37 | SNP 35 | SNP 34 | SNP 32 | SNP 31 | SNP 30 | SNP 28 | SNP 27 | SNP 26 | SNP 24 | SNP 23 | SNP 22 | SNP 20 | SNP 19 | SNP 16 | SNP 15 | SNP 14 | SNP 13 | SNP 12 | SNP 6 | SNP 5 | SNP 4 | SNP 3 | SNP 2 | SNP 1 | p-value | Aff % | Ctrl % | RR |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|--------|-------|-------|-------|-------|-------|-------|---------|-------|--------|------|
| T | A | A | C | C | A | C | G | A | A | C | T | T | A | T | T | G | A | A | T | T | T | G | A | A | 0.0006 | 22.2 | 15.4 | 1.58 |
| G | A | A | C | C | A | C | G | A | A | T | C | C | G | C | C | G | A | G | C | A | T | C | A | A | 0.533 | 2.2 | 2.8 | 0.78 |
| G | A | A | C | C | A | C | G | A | A | T | C | C | G | C | C | G | A | G | T | T | T | G | A | A | 0.24 | 3.0 | 2.0 | 1.52 |
| G | A | A | C | C | A | C | G | A | T | T | C | T | A | C | C | A | G | G | C | A | C | C | T | G | 0.302 | 2.6 | 1.8 | 1.46 |
| G | G | C | T | T | C | C | G | A | A | C | T | T | A | T | T | G | A | A | T | T | T | G | A | A | 0.258 | 2.0 | 3.0 | 0.68 |
| G | G | C | T | T | C | C | G | A | A | T | C | C | G | C | T | G | A | G | C | A | T | C | T | G | 0.0078 | 7.8 | 12.0 | 0.62 |
| G | G | C | T | T | C | G | A | G | T | T | C | T | A | C | C | A | G | G | C | A | C | C | T | G | 0.781 | 32.9 | 33.6 | 0.97 |

Allelic definitions and polymorphisms for SNPs in the two most significant haplotypes (in block B and C).

The analysis presented above represents a conservative analysis of the data since it restricted the analysis to SNPs with minor allelic frequencies of greater than 20%. To further understand the magnitude of the contribution of PDE4D to stroke in this 5 prime region, we repeated the analysis without such restrictions, including all SNPs selected for genotyping. We found a SNP haplotype for the two major subtypes of ischemic stroke, carotid and cardiogenic stroke (Table 3, Block C). This is a 5 SNP haplotype that covers an area of 48 kb and is just upstream of the 5'exon covering the presumed promoter region of isoform PDE4D7. It captures the same information as the 0 allele for marker AC008818-1. However, the SNP haplotype is more specific in the sense that it has a higher relative risk, *i.e.*, 2.3. This haplotype is carried by 47% of the patients and has the same population attributable risk (PAR) of 0.25. The polymorphisms and alleles for the SNPs are presented in Table 4A.

Table 4A

| SNP Name | Public name if available | Polymorphism | Position | Allele (nucleotide) |
|------------------------|--------------------------|--------------|----------|---------------------|
| SNP42 SNP5PDM361194 | rs153031 | A/G | 138806 | 0 (A) |
| SNP34 SNP5PDM368135 | rs27653 | C/A | 131865 | 0 (A) |
| SNP32 SNP5PDM370640 | rs456009 | C/T | 129361 | 1 (C) |
| SNP26 SNP5PDM379372 | rs40512 | G/A | 120628 | 0 (A) |
| SNP9 SNP5PDM408531 | (new) | G/A | 91470 | 0 (A) |

Table 4B

| Phenotype | SNP42 | SNP34 | SNP32 | SNP26 | SNP9 | p-value | # Affect | Aff. Freq.* | # Ctrl | Ctrl. Freq* | R-risk | PAR | info |
|-------------------------|-------|-------|-------|-------|------|----------|-------------|----------------|-----------|----------------|--------|------|-------|
| All stroke | A | A | C | A | A | 2.17E-05 | 988 | 0.19 | 652 | 0.12 | 1.8 | 0.16 | 0.604 |
| Cardiogenic/ carotid | A | A | C | A | A | 3.37E-07 | 313 | 0.236 | 652 | 0.119 | 2.3 | 0.25 | 0.616 |

• allelic frequency

5

The sequences for the microsatellite markers are as follows:

AC008879-2 amplimer:

ACAAAGAGCACCTTTCCAGTGGACAACCTAACTAAAGTGGTGTGATTTTGG
10 TATAAGTTTGTATGTGTA
TACATTTAGTTTTATTGTAACAAAGCAACTTGTACTTTTCACGTTTAAAA
(SEQ ID NO: 85)

* AC008879-2, allele 0 is the same allele as the minimum allele observed in
CEPH 1347-02, family 137, individual 02.

15

In summary, this single SNP haplotype (which is only one haplotype of the
several found above but is probably the most tightly associated to stroke) more than
20 doubles an individual's risk for cardiogenic and carotid stroke and accounts for 25%
of such strokes in Iceland. The other haplotypes described above provide additional
risk for stroke. The magnitude of this risk haplotype is comparable or higher than
the well-known clinical risk factors for stroke such as hypertension, diabetes,
hyperlipidemia, and smoking.

These SNPs show strong association in patients with cardioembolic and large vessel disease.

Table 5 and Table 6 show previously known microsatellite markers and novel microsatellites in sequence. Forward and reverse primers are shown.

Table 5 Previously Known microsatellite markers in sequence

| | Accession number | Forward primer | SEQ ID NO. | Reverse primer | SEQ ID NO. |
|---------|------------------|---------------------------|------------|----------------------------|------------|
| D5S2107 | GDB:614475 | AGCCTTTGGGCAACA | 15 | CAAAACCAACAGGAGTATGTACTTTT | 16 |
| D5S468 | GDB:593646 | AAATGAATGGTAGATTTAACCTGAG | 17 | TGGGAAAATAAATACATGCG | 18 |
| D5S2000 | GDB:608769 | TTATACCAGGAGAGTAGACTTTTTT | 19 | CATGCTAATTTCAAAATATGAGAG | 20 |
| D5S2091 | GDB:613806 | GCAATTTGTCATGTGCCA | 21 | GGTATTTCAATTCACAGCCAGTC | 22 |
| D5S2500 | GDB:683034 | TTAAAGGAGTGATCTCCCCC | 23 | GTACAGTACCTATGATCATGCC | 24 |
| D5S2080 | GDB:613188 | GCACTGTGAATTTCAAATG | 25 | GTCAGGGGACTGGGAT | 26 |
| D5S2018 | GDB:609957 | CCTGTAAACAAATGAAACCCACTGA | 27 | AGACTATGCTGTGTGTGTCCTG | 28 |
| D5S2071 | GDB:612756 | TCTGGGTTTACAAACCTTCAA | 29 | TAACTGGCTTGGCCCG | 30 |

| Table 6 Novel microsatellites in sequence: | | SEQ ID NO. | Reverse primer | SEQ ID NO. |
|--|---------------------------|------------|---------------------------|------------|
| | Forward primer | | | |
| DG5S382 | CAGTAAATAGTTGCTTCAGGCATT | 31 | CTCATACTCTGCGTGGCTTG | 32 |
| AC008829-5 | AGGGCTAAGTGGATCACAGC | 33 | AGAGGTCTTTGCCACTGTGT | 34 |
| AC008833-2 | TCTGCAAGACTCTCGGTGCT | 35 | TGCAGATCTCATATTTCCATGTTT | 36 |
| AC008833-3 | TCTGCCCTTTGTTCCCTCATC | 37 | GTCAAAGGAGTGATGGCAGT | 38 |
| AC022125-3 | AAATGACTGCCTCCACAA | 39 | GGGAAATCATACTGCCCTCA | 40 |
| AC008833-6 | AAACATAGCCACCCTGTTC | 41 | TCCAAAGCCCTTAGCTTAATCA | 42 |
| D17-C | GCTCCCTGGACTGTGGTAA | 43 | GCCACATTGCTGTACACATTT | 44 |
| D17-B | TTTTCAGGGCTGGGTAGAA | 45 | TCCAAAGGAAGTGAAATCAGTG | 46 |
| D17-D | CTAACCCTATCCTCACCCCAAT | 47 | TGTGGCATACAGGGAAAGTGA | 48 |
| AC008804-1 | GTGCTGGAAATTTGGCTCCTA | 49 | CAAAACATCATTTTGCCTTGC | 50 |
| AC008804-2 | TCCCAACGATAGCTGTTC | 51 | GAATTAGGACGGTGGCTCAA | 52 |
| AC008804-3 | TTTGCAATTCATCACTCAATCG | 53 | CCCGTAGCATCTGATCCAGT | 54 |
| D17-H | AGAAAGCTTCCCTCCACTG | 55 | CATTCCAGCCTGAGCTACAA | 56 |
| D17-G | TGGGCTCCAAATTAATCCTCC | 57 | TGCAGTTTGCACTCTCCTTG | 58 |
| AC027322-12 | TTATCTGTTCCTCCATGCTTTT | 59 | TGTTACATCTTGATCTATGACGTTT | 60 |
| AC027322-10 | TGTAATCCTGCATCCCTTGTT | 61 | GGAATAACCCAAAGTAATTTAGTGA | 62 |
| AC027322-9 | TCGTGCCAAGATGAAAATGA | 63 | AAACCTCCCTGATCATCTGAA | 64 |
| AC027322-8 | ACAGAGGAGCAAAGGAATCA | 65 | TTGGCAGCAATCACTCTCTG | 66 |
| AC027322-3 | CCCCATTTGGATGATGGTAA | 67 | TGAGAACATCTAAACGCTTTTTCAA | 68 |
| AC027322-5 | GGCACAGATAACTGGGAAGC | 69 | CCCCCAAAGTACTGCATAAA | 70 |
| DG5S397 | ATGTTGGCATTTGGTGAGGT | 71 | CACCTGTCCCTTTGGAGGTA | 72 |
| AC008879-2 | TTTAAACGTGAAAAGTACAAATTGC | 73 | ACAAAGAGCACCTTTCCAGTG | 74 |
| *AC008818-1 | TGCTTGGTGAAAGGAATAGCC | 75 | GAGCCTGGGTTCTCAGGAAT | 76 |
| **AC008879-3 | GGCAAGAACAGTTTGGAGGA | 77 | GACTGCTGTTTGCTGGTTGA | 78 |
| AC020733-1 | AAATGGCTATAAAGTGTCTTTGAAC | 79 | CGGTCTCAACAACCAAGAACAA | 80 |
| AC016591-2 | CAGAAACACACAGAAAGTCATTCAA | 81 | CAGACCCCAATTAATGGCAAAA | 82 |
| DG5S405 | TCTGTCTTCTTTGACCCCATGAAT | 83 | CAACACAGCGAGACCTCATC | 84 |

* Product Size 194, tetranucleotide repeat

** Product Size 150; dinucleotide repeat

*

**

Table 7
Correlation between at-risk alleles for markers AC008818-1, SNP45 and SNP41.
Estimates of LD (correlation) between the at-risk alleles, allele 0 for marker AC008818-1, allele
G for SNP45 and allele A for SNP41, the three most significant disease associated genetic
5 markers.

A. Combined cardiogenic and carotid patients

| | | R^2 | | |
|----|------------|-----------|------------|--------------------|
| | | Frequency | AC008818-1 | |
| D' | AC008818-1 | 0.355 | | SNP 45 0.090 0.076 |
| | SNP 45 | 0.863 | 1 | 0.943 |
| | SNP 41 | 0.860 | 0.906 | 0.981 |

B. Controls

| | | R^2 | | |
|----|------------|-----------|------------|--------------------|
| | | Frequency | AC008818-1 | |
| D' | AC008818-1 | 0.255 | | SNP 45 0.091 0.078 |
| | SNP 45 | 0.780 | 1 | 0.920 |
| | SNP 41 | 0.768 | 0.924 | 0.968 |

Table 8

Association of risk factors.

5 Association of microsatellite AC008818-1 at-risk allele 0, SNP45 allele A and haplotype G0H_C respectively with various risk factors.

Cases are stroke patients with risk factors and controls are stroke patients without the risk factors. P-values are two-sided.

| | AC008818-1 : Allele 0 | | | | | SNP45 : Allele A | | | | | Haplotype G0H _C | | | | |
|-----------------------------|------------------------|-------|---------------------------|-------|---------|------------------------|-------|---------------------------|-------|---------|----------------------------|-------|---------------------------|-------|---------|
| | Cases with risk factor | | Cases without risk factor | | P-value | Cases with risk factor | | Cases without risk factor | | P-value | Cases with risk factor | | Cases without risk factor | | P-value |
| | N | Frq. | N | Frq. | | N | Frq. | N | Frq. | | N | Frq. | N | Frq. | |
| Hypertension | 477 | 0.303 | 203 | 0.303 | 1.000 | 416 | 0.172 | 181 | 0.188 | 0.510 | 503 | 0.134 | 216 | 0.123 | 0.634 |
| Hyper-cholesterolemia | 274 | 0.336 | 312 | 0.271 | 0.025 | 242 | 0.216 | 277 | 0.186 | 0.516 | 287 | 0.153 | 329 | 0.104 | 0.026 |
| Diabetes | 93 | 0.274 | 424 | 0.310 | 0.379 | 79 | 0.196 | 398 | 0.176 | 0.422 | 100 | 0.127 | 455 | 0.133 | 0.857 |
| Peripheral artery occlusive | 133 | 0.297 | 357 | 0.305 | 0.815 | 116 | 0.181 | 340 | 0.176 | 0.921 | 138 | 0.121 | 388 | 0.132 | 0.697 |
| Coronary artery disease | 179 | 0.302 | 429 | 0.318 | 0.588 | 153 | 0.170 | 406 | 0.182 | 0.662 | 181 | 0.122 | 467 | 0.141 | 0.444 |
| Early onset (< 68) | 349 | 0.294 | 462 | 0.304 | 0.137 | 314 | 0.186 | 430 | 0.173 | 0.538 | 380 | 0.128 | 506 | 0.125 | 0.876 |
| Males vs females | 457 | 0.291 | 358 | 0.310 | 0.414 | 420 | 0.181 | 303 | 0.168 | 0.575 | 489 | 0.122 | 370 | 0.141 | 0.315 |

Discussion of Stroke Gene Identification

10 Genealogy, a comprehensive population-based list of broadly defined stroke patients and non-parametric allele sharing methods have been combined to successfully map a major gene to chromosome 5 for one of the most complex diseases known. We then used a large case-control association study that showed that PDE4D is the gene in this location that is the gene conferring substantial risk for stroke. This is the first gene
15 ever mapped and isolated for the common forms of stroke. There was no correlation

between the contribution of the families to this gene location and hypertension, diabetes or hyperlipidemias and this gene does not match any known gene contributing to these risk factors. The types of stroke studied in this work do not reflect a rare or Icelandic-specific form of stroke; rather, the diversity of the stroke phenotypes in Icelanders as well as risk factors are similar to those of most other Caucasian populations (Agnarsson, U., *et al.*, *Ann. Intern. Med.*, 130:987 (1999); Eliasson, J.H., *et al.*, *Læknablaðið*, 85:517-25 (1999); Sveinbjörnsdóttir, S., *et al.*, Systematic registration of patients with Stroke and TIA admitted to The National University Hospital, Reykjavik, Iceland, in 1997, XIII. Meeting of the Icelandic Association in Internal Medicine, Akureyri, Iceland (Valdimarsson, E.M., *et al.*, *Læknabladid* 84:921 (1998)).

The magnitude of the risk and the frequency of the disease haplotypes in the general population confirm that we have mapped a gene for the common forms of stroke and not some rare form of stroke. This gene almost doubles one's risk for stroke in general, and more than doubles one's risk for the two most common subtypes of stroke, carotid and cardiogenic stroke. In addition, the most common disease haplotype has a population attributed risk of 25% (which means it accounts for 25% of the patients) and there are other haplotypes that we describe herein that are less common that accounts for other patients. Thus PDE4D is a major cause of stroke and its relative risk rivals those of hypertension, smoking, diabetes, and hyperlipidemia. PDE4D shows tighter correlation to the forms of stroke dependent on atherosclerosis (carotid and cardiogenic stroke) and it is expressed in cell types known to be important for atherosclerosis such as vascular smooth muscle cells, macrophages, and endothelial cells. This suggests that the strong effect that PDE4D variation has on stroke risk is through its role in the vascular biology of atherosclerosis (see discussion at the end of the examples). Example 2 details our sequencing of the entire PDE4D gene and the definition of its exon-intron structure based on new and old cDNAs, and Example 3 shows that the expression pattern of PDE4D isoforms correlates with a stroke associated haplotype.

EXAMPLE 2: SEQUENCING AND CHARACTERIZATION OF THE HUMAN GENE AND ITS RNA/PROTEIN ISOFORMS

Sequence of the Stroke Gene Region

5 At the start of our work, there was little genomic sequence available in the public domain covering the stroke gene region. Therefore, we sequenced approximately 3 Mb of the area defined by one drop in lod. The locus on 5q12 indicated in the genome wide scan was physically mapped using bacterial artificial chromosomes (BACs). A set of overlapping clones for a 20 cM region was assembled through a combination of

10 hybridization and BAC-fingerprint walking. Eighteen BACs (bacterial artificial clones): (RP11-164A5, RP11-188I15, RP11-313P15, RP11-631M6, RP11-103A15, RP11-489L13, RP11-621C19, RP11-113C1, RP11-567M18, RP11-412M9, RP11-151G2, RP11-151F7, RP11-281M3, RP11-421L6, RP11-1A7, RP11-68E13, RP11-379P8, and RP11-422K3) covering the minimum tiling path of the one LOD interval were analysed

15 using shotgun cloning and sequencing. Dye terminator (ABI PRISM BigDye) chemistry was used for fluorescent automated DNA sequencing. ABI prism 377 sequences were used to collect data and the Phred/Phrap/Consed software package in combination with the Polyphred software were used to assemble sequences (See Table 9A and 9B)

20 Publicly available sequences (AC008836, AC073546, AC021603, AC008498, AC016435, AC021601, AC016591, AC008818, AC008879, AC008934, AC011929, AC027322, AC008111, AC020924, AC026693, AC012315, AC08804, AC008791, AC020975, AC008833, AC008829, AC022125, AC008790, AC026095, AC066693, AC008852, AC016642, AC034250, AC025179, AC08814, AC008926, AC010391, AC016635 and AC016604) from this region were assembled with the obtained sequence

25 and a 3.7 Mb sequence (with 22 gaps) was generated. Comparison of the current public human assembly (NCBI BUILD 33) to our sequence of the STRK1 locus only showed a minor discrepancy.

The BAC clones we sequenced are from the RCPI-11 Human BAC library (Pieter deJong, Roswell Park). The vector used was pBACe3.6. The clones were picked

into a 94 well microtiter plate containing LB/chloramphenicol (25 µg/ml)/glycerol (7.5%) and stored at -80°C after a single colony has been positively identified through sequencing. The clones can then be streaked out on a LB agar plate with the appropriate antibiotic, chloramphenicol (25 µg/ml)/sucrose (5%).

5

Table 9A

| Sequenced at Decode (BAC name) | Comment | Accession number |
|-----------------------------------|---------|------------------|
| RP11-621C19 | 1 | AC020733 |
| RP11-113C1 | 2 | |
| RP11-412M9 | 2 | |
| RP11-151G2 | 2 | |
| RP11-151F7 | 2 | |
| RP11-281M3 | 2 | |
| RP11-421L6 | 2 | |
| RP11-68E13 | 2 | |
| RP11-379P8 | 2 | |
| RP11-1A7 | 1 | AC008111 |
| RP11-422K3 | 2 | |

Key to "Comment" column:

1= This BAC has a publicly available sequence,

it was sequenced at Decode to make sure the sequence was correct

2= Only BAC end-sequence available for this BAC publicly.

10

Table 9B

| Sequences available from GenBank (BAC name) | Accession number | Status of sequence |
|--|-------------------------|---------------------------|
| RP11-621C19 | AC020733 | 17 unordered pieces |
| CTD-2003D5 | AC016591 | complete sequence |
| CTD-2210C1 | AC008879 | 7 unordered pieces |
| CTD-2124H11 | AC008818 | complete sequence |
| CTD-2301A11 | AC008934 | complete sequence |
| RP11-16B11 | AC011929 | 7 unordered pieces |
| CTC-261E10 | AC026693 | complete sequence |
| CTD-2027G10 | AC027322 | complete sequence |
| RP11-1A7 | AC008111 | 8 unordered pieces |
| CTD-2122K7 | AC012315 | complete sequence |
| CTD-2085F10 | AC008804 | complete sequence |
| CTD-2040J22 | AC008791 | complete sequence |
| RP11-235N16 | AC020975 | 16 ordered pieces |
| CTD-2146O16 | AC008833 | complete sequence |
| CTD-2084I4 | AC022125 | 17 ordered pieces |
| CTD-2140K22 | AC008829 | 26 ordered pieces |
| CTD-2124D11 | AC020924 | 7 ordered pieces |
| RP11-731H6 | AC026095 | 21 unordered pieces |

PDE4D Gene; Identification of New Exons and Splice Variants

The gene, human cAMP specific phosphodiesterase 4D (HPDE4D) was identified in the sequenced region by BLAST of our novel genomic sequence with the cDNAs/EST databases from GenBank. In addition, we ran RT-PCR reactions and 5 prime and 3 prime RACE reactions using cDNA libraries generated from a variety of tissues including human aorta. The primer sites used corresponded to known or exons predicted from our genomic sequence using Genscan, and Fgene. We found several novel cDNAs and matched them to the 3Mb sequence in and around PDE4D. The genomic sequence covering all known and novel exons in PDE4D so far is approximately 1,550,000 bases in length.

We defined new alternative transcripts which together with previously known transcripts showed that the *PDE4D* gene contains 22 exons over at least 1.5 Mb and overlaps with the *PART1* gene whose transcript is on the other strand at the 5' end. The *PDE4D* gene has at least 7 promoters and encodes 8 protein isoforms. All isoforms have an identical C-terminal catalytic domain but differ at the N-terminal regulatory domain. Six of the 8 forms are so called long isoforms. Each of them have unique N-terminal regulatory domains but they are all characterized by two highly conserved regions found in all PDE4 subfamilies, *i.e.* upstream conserved regions 1 and 2 (UCR 1 and 2). The six long forms differ from each other by unique alternative 5 prime exons which predicts six alternative promoters that are each upstream of the corresponding 5 prime exon. The remaining two are the so-called short forms, variants that lack the UCR 1 (Houslay, M.D. & Adams, D.R., *Biochem J*, 370, 1-18 (2003)). The five previously known isoforms are encoded by 17 exons distributed over a segment of 0.9 Mb.

The three new exons D7A-1, D7A-2 and D7A-3 are spliced to one another and together splice onto exon LF1 forming the splice variant we named PDE4D7 (FIG. 3). Exon D7-1 is non-coding. Exons D8 and D9 are spliced by themselves onto exon LF1 forming two splice variants we named PDE4D8 and PDE4D9, respectively (FIG. 3).

In terms of genomic structure, the D7A exon extends the 5' end of PDE4D by 590,000 bp, and the D8 and D9 exons lie between exons D3 and LF1 (physical position of exons presented in Table 2C). The new PDE4D7 isoform has an open reading frame extending into LF1, resulting in additional 91 amino acids at the N-terminus of the predicted protein. The D8 and D9 5' exons contain a long 5' UTR, followed by an ATG near the end of the exons that extends an ORF into LF1 resulting in a novel N-terminal segments of 22 and 30 amino acids in the PDE4D8 and PDE4D9 predicted proteins, respectively. The new splice variants were verified by RT-PCR on different cDNA tissue panels and subsequent cloning and sequencing of the products.

The *PDE4D* gene encodes at least eight different isoforms. Six of the eight forms are the so-called long isoforms. Each of them has a unique N-terminal regulatory domain but they are all characterized by two highly conserved regions found in all PDE4 subfamilies, *i.e.*, upstream conserved regions 1 and 2 (UCR 1 and 2). The remaining two isoforms are the short forms, variants which lack the UCR 1.

Three PDE4D isoforms have been submitted to GenBank by Memory Pharmaceuticals on September 16, 2002 and December 17, 2002, under accession numbers AF536975 (isoform named PDE4D6), AF536976 (named PDE4D7) and AF536977 (named PDE4D8). See also PCT WO 01/00851, published January 4, 2001. The sequence AF536977 corresponds to our earlier reported PDE4D6 isoform and AF536976 corresponds partly to our earlier reported PDE4D7 isoform, however the first untranslated exon we named D7-1 is missing from this sequence. The sequence AF536975 is a new short PDE4D isoform. We have therefore changed the isoform names accordingly herein as follows: PDE4D6 is now called PDE4D8, PDE4D7 is now called PDE4D7 and PDE4D8 is now called PDE4D9. We have submitted the new PDE4D splice variants, PDE4D7 and PDE4D9 to GenBank (Accession numbers AY245866 and AY245867, respectively).

We have in addition identified 17 putative exons upstream of LF1, based on ESTs, mouse homologies and GeneMiner exon predictions. Primers designed from these exons were used in conjunction with primers from LF1 and exon3 for RT PCR in the hope of identifying novel exons. Novel exons were in turn used to design primers for various RT-PCR reactions. We also used 5'RACE primers, designed from the known exons upstream of LF1. We have to date identified 14 new exons, including exons belonging to UniGene Cluster Hs.343602 that have now been connected to LF1.

For the 5' RACE reactions we used cDNA made from heart, SkNAS (neuroblastoma cell line) and HVAEnd 5050 (endothelial cell line). For RT-PCR reactions a number of cDNAs made of total RNA were used (see below)

Novel exons in Table 10A are in italics; previously know PDE4D exons in white. Exon 3 of EST AW272330 is included on the table as a representative of the 3' of ESTs from UniGene cluster Hs 343602. The positions given are from SEQ ID NO: 1. Note the different splicing of 4D9-3.2, 4D9-3.1, and AW272330exon3.

Total RNA was isolated from HeLa, SkNAs and Jurkat 77 cell cultures according to manual, using the TRIZOL[®] reagent provided by GibcoBRL. We used the GeneRacer[™], ThermoZyme[™] and TOPO TA cloning[®] (containing pCR[®]2.1-TOPO[®]) kits from Invitrogen following the manufacturer's protocol.

Table 10A

| exon # | EXON | SEQ ID 1 | SEQ ID 1 | supported by EST(s) |
|--------|---------------|----------|----------|---------------------|
| 1 | 4D7-A | 108127 | 108217 | Yes |
| 2 | PDE4D7-1 | 142207 | 142328 | PDE4D |
| 3 | 4D7-4 | 257650 | 257705 | Yes |
| 4 | 4D7-8 | 288224 | 288393 | No |
| 5 | 4D7-B | 295203 | 295251 | No |
| 6 | 4D7-5 | 352169 | 352317 | No |
| 7 | 4D7-6 | 441914 | 442036 | No |
| 8 | PDE4D7-2 | 444645 | 444775 | PDE4D |
| 9 | 4D7-C | 482438 | 482719 | No |
| 10 | 4D7-7 | 597399 | 597534 | Yes |
| 11 | 4D7-9 | 626020 | 626092 | No |
| 12 | PDE4D7-3 | 641649 | 641878 | PDE4D |
| 13 | PDE4D4 | 736254 | 737226 | PDE4D |
| 14 | PDE4D5 | 861791 | 862202 | PDE4D |
| 15 | PDE4D3 | 1044051 | 1044190 | PDE4D |
| 16 | 4D9-1 | 1069544 | 1069629 | Yes |
| 17 | 4D9-2 | 1069936 | 1069993 | No |
| 18 | 4D9-3.2 | 1071661 | 1071795 | Yes |
| | 4D9-3.1 | 1071668 | 1071795 | |
| | AW272330exon3 | 1071668 | 1071901 | |
| 19 | 4D9-4 | 1121821 | 1121892 | No |
| 20 | 4D9-5 | 1247621 | 1247696 | No |
| 21 | PDE4D8 | 1273404 | 1273709 | PDE4D |
| 22 | PDE4D9 | 1354347 | 1355128 | PDE4D |
| | LF1exon | 1414511 | 1414702 | PDE4D |

5 *Sequence of New Exons:*

>4D7-A
GGCCTCGAGCAGAACTTCCCATTGAGTGGGACCAAGAAGAGCATACAAAG
CTGAAATGTTCTCCAGAAGTTGATTTC CAATGGGGATAAA (SEQ ID NO: 88)

10 >4D7-4_ From Forward Primer
TGATTACAGGTTTTAGAGAAGAGGAACAATGCTTCCTCTGAGCCTGAAGAA
AAGAA (SEQ ID NO: 89)

>4D7-8

AGTTCTGACCATGTCCTGTGTCACTCTCAAGCAGAGATTGAAAATGACATTC
GTCCTTTACTTGTTCCAAGGAAGCAAACATTTTATAGTTTGAAACTGTTTCTC
TTGCATTTGCTTTGCAAGAGGTTTGCAGAAGTTAAGCCTCATGGAGTCTTCTC

5 TCCTTAACTTAA (SEQ ID NO: 90)

>4D7-B

TGTGAAGAATTTGGAAATTGCAAGGAGCATGGGAAGGAGATGATTTGGG
(SEQ ID NO: 91)

>4D7-5

10 GAATGAAGAGGAAATCAAGACATACTTAGATAAAAAACAGATTATCACCAGG
AGATCTGCTGTAAAAGAATGGCTAAAGGAAGTTAGCTAAGCAGAAAGGAAG
TAACATAAAAAGGAACCTTGGAACATCAGGGAGGACAAAAGAACATG (SEQ
ID NO: 92)

>4D7-C

15 TTTCTCTTTCTCCAATCACTCACTCTGGAGGCAGCTAGCTGTCAACTCACAAA
GACACTCAAGCAGCCTATGGGAAGAAGGCCACATGGTAAAATATGGAGGCCT
CCAGCCAACAGTCAGCAAGGAAGTGAAGCAAGTCAACAACCATGTGAGTGA
CTCGAGAAGTGCTTCTCTAGCTCCAGTTGAGACTTGCAGTAGCAGCAGCCTC
AGCTGGCGGCTTGACTGCAATCTCTTGAGAGACCCTAAGCTCTCCTGAATTC
20 TTGATCCTTAGAAACTGTGTGAG (SEQ ID NO: 93)

>4D7-6

GGTCTAGCTGTGTCCCAGAGAGCAACTTCCCTTTTCAAGGCAGCCCCTCTG
TGTGATGCTTTTTCCTAGGTATGGGCAACCCATCCCTCCTAGGGTGAAAAC
TCGCTGTTGCTAGTTCCAG (SEQ ID NO: 94)

25 >4D7-7

AATGATGCCGTATTATTCTCCTGACCTAACTTCAAAGAAATAAAGAGTTTGC
AAGAAGAACTGCAGTTCTTCAAAGTACGCAATATGGATTTCCAAGATGAAT
GTAGTTTCTCTCTCTGAGGAATTCTGAACAGTG (SEQ ID NO: 95)

>4D7-9

30 GACTTGAGCATCTGAAGATTTTGGTTTCTGCAGAGGGTGGGAAAGGTTGAAC
CAATCCCCCATGGATACCAAG (SEQ ID NO: 96)

>4D9-1

GGCTTTCCAGATCCCTGAAGATAAAATACAAACTCTCCAACAAGACCTTT
TGGCCATCAGGAACGCAGCACCTGGCTCTCTCACTA (SEQ ID NO: 97)

35 >4D9-2

AAAGTCGCAGAGATAGCGGAGAACAAGAACCAGATCTCACAGTCATGGTGC
CAAAAGA (SEQ ID NO: 98)

>4D9-3.1

40 CTGTTACCCTAGCATGACTGCTTCAGCGAAGAGATAAGAGCTTCTTTGACTT
TTTCCACTGGAATTTTTCATGCCAGAAGAAATTGAACATGTGAGCCTGGTGT
CTGGAAGAGTAGCCTGGATTTATG (SEQ ID NO: 99)

>4D9-3.2

AATTCAGCTGTTACCCTAGCATGACTGCTTCAGCGAAGAGATAAGAGCTTCT
TTGACTTTTTTCCACTGGAATTTTTCATGCCAGAAGAAATTGAACATGTGAGC
CTGGTGTCTGGAAGAGTAGCCTGGATTTATG (SEQ ID NO: 100)

>4D9-4

5 TTCCTTGATAGTTCCAATATCTGTAATCTTGTTGGTCTACCTGTGCAGTTTAT
TCCACTGATTGTCTCTCAG (SEQ ID NO: 101)

>4D9-5

GCGAAAATACTGAGGCTCAACAGACATAAAATGGCTTGAGTTACCAGGCTA
CAGTAGAACTAGGATTTTCAGTCCAG (SEQ ID NO: 102)

10

Splicing of the Exons as identified by RT-PCR/RACE

New exons are in italics.

15 RT4D7: 4D7-1 + 4D7-2 + 4D7-3 + LF1
RT1: 4D7-1 + *4D7-8* + 4D7-2 + 4D7-3 + LF1
RT2: *4D7-4* + 4D7-2 + *4D7-9* + 4D7-3 + LF1
RT3: 4D7-2 + 4D7-3 + LF1
RT4: 4D7-1 + 4D7-2 + *4D7-7* + 4D7-3 + *4D9-1* + *4D9-3.1*
20 RT5: 4D7-1 + 4D7-2 + 4D7-3 + *4D9-2* + *4D9-3.1*
Race6: *4D7-A* + *4D7-B* + 4D7-2 + *4D7-C* + 4D7-3
RT7: 4D7-1 + *4D7-4*
RT8: 4D7-1 + *4D7-5* + 4D7-2
RT9: 4D7-1 + *4D7-6* + 4D7-2
25 RT10: *4D9-1* + *4D9-3.1* + LF1
RT11: *4D9-2* + *4D9-3.2* + LF1
RT12: *4D9-3* + *4D9-4* + LF1
RT13: *4D9-3* + *4D9-4* + *4D9-5* + LF1
RT14: *4D9-3* + LF1
30

Detection of variants in cDNA from various tissues:

Table 10B

| | RT4D7 | RT1 | RT2 | RT3 | RT4 | RT5 | RT6 | RT7 | RT8 | RT9 | RT10 | RT11 | RT12 | RT13 | RT14 |
|-----------------|-------|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|------|------|------|------|
| Bone Marrow | - | - | - | * | - | - | n | n | n | n | n | - | - | - | - |
| Brain | + | - | - | * | - | - | n | n | n | n | n | n | - | + | - |
| fetal Brain | + | - | - | * | - | - | n | n | n | n | n | n | - | - | - |
| colon | + | + | - | * | - | - | n | n | n | n | n | n | - | - | - |
| Heart | - | - | - | - | - | - | - | - | - | - | - | - | - | - | + |
| HVAEend 5050 | - | - | n | n | n | n | + | - | - | - | + | + | n | n | + |
| Kidney | + | - | + | + | - | - | n | n | n | n | n | n | - | - | - |
| Liver | - | - | - | * | - | - | n | n | n | n | n | - | - | - | - |
| Placenta | - | - | - | * | - | - | n | n | n | n | n | - | - | - | - |
| Prostate | + | - | - | * | + | - | n | n | n | n | n | n | - | - | * |
| Salivary gland | + | - | - | * | - | - | n | n | n | n | n | n | - | - | - |
| Skeletal Muscle | - | - | - | - | - | - | n | n | n | n | n | n | + | - | + |
| SkNAS cell line | + | - | n | * | n | n | - | + | + | + | - | - | n | n | - |
| Spinal Cord | - | - | - | - | - | - | n | n | n | n | n | n | - | - | * |
| Spleen | - | - | - | * | - | - | n | n | n | n | n | n | - | - | + |
| Testis | - | - | - | - | - | - | n | n | n | n | n | n | - | - | * |
| Thymus | + | - | - | * | - | - | n | n | n | n | n | n | - | - | - |
| Thyroid | + | - | - | * | - | + | n | n | n | n | n | n | - | - | + |
| Trachea | + | - | - | * | - | - | n | n | n | n | n | n | - | - | - |
| Uterus | - | - | - | - | - | - | n | n | n | n | n | n | - | - | + |
| other# | - | - | - | - | - | - | n | n | n | n | n | n | - | - | - |

+ = present and verified by sequencing

5 * = product of the correct size present; not yet verified by sequencing

- = not detected

n = not checked

These are: Adrenal Gland, Fetal Liver, Cerebellum, Lung, Small Intestine.

10 Two of the variants that are more widely expressed appear to be mutually exclusive: 4D7 [with 4D7-1 as first exon] was detected in 10 cDNAs while RT14 is found in 9cDNAs. Of these thyroid and prostate are the only tissues common to both variants.

The 13 new RT and RACE variants presented above (we had previously described the 4D7 variant), do not add any new translated sequence. The RT1 product is expected to be the same as the 4D7 putative protein. In variants RT2 and Race6 the exons between 4D7-2 and 4D7-3 interfere with the ORF with the first AUG and ORF being just inside LF1. Similarly Exon 4D9-3 contains stop codons in all 3 reading frames and Variants RT10, 11,12,13, and 14 having their ATG initiation codon inside LF1. It is not clear whether variants RT4 and RT5, which contain exon 4D9-3 extend to LF1 or have their 3' at 4D9-3 (the latter possibility is supported by the EST data).

It is noteworthy that all variants except 4D7, RT3 and RT14 have been observed only in one of the cDNAs. Although all the new exons (except 4D9-3.1) have an AG/GT splice signal, it is plausible that these variants represent rare or aberrant events with little physiological significance.

The following exons contain Alu repetitive element sequences: 4D7-5 and 4D7-C. The gene specific reverse (3') primer was designed for PDE4D exon LF1 (5' GGCAATGGAGGAGTTCCGGGACA TA-3'; SEQ ID NO: 87 origin from Homo sapiens).

A contig for the incomplete genomic sequence of the PDE4D gene was submitted by others in November 2000 (GenBank entry NT_023193 by International Human Genome Project collaborators). The size of the contig is 614 481 bp (including gaps) whereas our novel genomic sequence for the whole PDE4D region (*i.e.*, from the first exon for PDE4D variant) is close to 1,690,000 bp and contains no gaps. The contig NT_023193 comprises only 11 exons of the PDE4D gene (in FIG. 3, exons 4D1/D2 - 11) and the 5' differently spliced exons are missing in the contig (in FIG. 3, exons D4, D5, D3, D8, D9, D7A-1, D7A-2, D7A-3, LF1, LF2, LF3 and LF4).

Table 13: New Isoforms

| Isoform Name | | | | Cell line |
|---------------------|------|----------|--------|-----------|
| | Exon | | Size | |
| PDE4D7 | D7-1 | 5' | 122 bp | SKNAS |
| PDE4D7 | D7-2 | Internal | 131bp | SKNAS |
| PDE4D7 | D7-3 | Internal | 230 bp | SKNAS |
| PDE4D9 ¹ | D9 | 5' | 782 bp | HeLa |

5 ¹ Formerly referred to in previous applications as PDE4D8

The sequences are as follows:

D7A-1:

10 ATAGTTGGCGTACCCTGAGGCCTGCCAGTTCCTGCCTTAATGCATATGTAGT
CGTAATTGAGTTCTGACACGGCCTTGGATGTTTCTGTCCTAAATAGCTGACA
TTGCATCTTCAAGACTGT

D7A-2:

15 CATTCCAGTTGGCTTTTGAGTGGATACGTGCAGTGAGATCATTGACACTGGA
AACACTAGTTCCCATTTTAATTACTTAAAACACCACGATGAAAAGAAATACC
TGTGATTTGCTTTCTCGGAGCAAAAGT

D7A-3:

20 GCCTCTGAGGAAACACTACATTCCAGTAATGAAGAGGAAGACCCTTTCCGC
GGAATGGAACCCTATCTTGTCCGGAGACTTTCATGTCGCAATATTCAGCTTC
CCCCTCTCGCCTTCAGACAGTTGGAACAAGCTGACTTGAAAAGTGAATCAGA
GAACATTCAACGACCAACCAGCCTCCCCCTGAAGATTCTGCCGCTGATTGCT
ATCACTTCTGCAGAATCCAGTGG (SEQ ID NO: 11; includes D7A-1, D7A-2 and
25 D7A-3)

New predicted amino-terminal protein sequence from above (PDE4D7):

30 MKRNTCDLLSRSKSASEETLHSSNEEEDPFRGMEPYLVRRLSCRNIQLPPLAFRQ
LEQADLKSESENIQRPTSLPLKILPLIAITSAESS (90 amino acids) (SEQ ID NO: 12)

D9:

35 TTCTCACTGCCCTGCGGTGTTTTGAACTGCCTTCTTACAGACGTCATACAGCC
CTTGAGGAATAGTTTCTGCCTGGTGAGATTGAATGATAGTTCTCATTCAAA
AACCCTGGATTCTAAGCAGGGACACACAGAAATTACTTTCGCAGGTAAATC
AGCCCACCCAGCCAAAGTGTGGAGAGATTTGTTTCCTTGGCTGACTTCTTTGC

TCCACGGAGAGGAGTGTTTTCTGTGCTTGCCCTGAAATGGAACTTCCTTGA
CAGCTCTCCCGTGTTACAGTACCTCCCGGTCATTTTCTTTTTCTCTCTCTAC
CTGCGCTCTTCGAGTGTGAGAAACCTTTAAAGCTGTTACTATGGAATTGCAA
AAAAGAGATCAAGTGACTCTTTCCTATGCTGGTTTCCCTTGTGACCCAGAT
5 GAAGAATCAATTCAGAATTCAGTTCCTCCCTTGGCATTGCAAGACACAGAAG
AACTGTCACTTCCTAACAGCCTAGTACTGGAGTAAATTCAGTATGAAGGAA
GAAAGCGCTCCTGCGTGTTAGAACCTTGCCCATGAGCTGGACCGAGGACAG
GAGATGGACTCCAGGAAAATTGGATTTCTTCAAGCAGCCTCCCTTGGAAATG
GAATATCTTTAAAATCTTCTTTGCAGAAAGACAGTTAGAATGTATTAATCAG
10 AATAGTTGAAGACTTATTTTCCTTTTTTATTTTTTTTCAAAATGAGCATTATTAT
GAAGCCAAGATCCCGATCTACAAGTTCCTAAGGACTGCAGAGGCAGTTTG
(SEQ ID NO: 13)

New predicted amino-terminal protein sequence from above (PDE4D9):
15 MSIIMKPRSRSTSSLRTAEAV (21 amino acids) (SEQ ID NO: 14).

20

Table 11

Publically Available SNPS; SNP ID No. from NCBI Database

| | | | | | |
|-----------|----------|-----------|-----------|-----------|-----------|
| rs286155 | rs27960 | rs27221 | rs149079 | rs789615 | rs37708 |
| rs286156 | rs27564 | rs27653 | rs149324 | rs401207 | rs37709 |
| rs2061250 | rs27565 | rs26955 | rs153067 | rs364917 | rs789389 |
| rs286150 | rs26948 | rs26956 | rs40354 | rs404202 | rs1423247 |
| rs206789 | rs40131 | rs153031 | rs26951 | rs440607 | rs874768 |
| rs1823062 | rs26949 | rs185190 | rs153029 | rs411255 | rs2042315 |
| rs1823063 | rs26950 | rs37762 | rs27223 | rs615429 | rs918590 |
| rs1445852 | rs26954 | rs37761 | rs27222 | rs789396 | rs918591 |
| rs766119 | rs26953 | rs1423471 | rs251726 | rs37684 | rs918592 |
| rs956721 | rs152324 | rs27224 | rs1862589 | rs1445893 | rs1115372 |
| rs248910 | rs35385 | rs1645013 | rs702556 | rs37685 | rs1345782 |
| rs248912 | rs40512 | rs1423472 | rs702554 | rs1086121 | rs1363862 |
| rs187481 | rs35386 | rs27220 | rs441391 | rs42222 | rs1423248 |
| rs153152 | rs35387 | rs1423473 | rs446883 | rs37707 | rs1423246 |

| | | | | | |
|-----------|-----------|-----------|-----------|-----------|-----------|
| rs1862614 | rs1995780 | rs1435077 | rs159624 | rs1008709 | rs298088 |
| rs2194256 | rs1508865 | rs1369287 | rs1159470 | rs1027747 | rs298087 |
| rs889305 | rs952110 | rs1017410 | rs159622 | rs869685 | rs1421401 |
| rs2113071 | rs1533019 | rs1017409 | rs256349 | rs869686 | rs298086 |
| rs2113072 | rs2117552 | rs1435076 | rs256348 | rs924880 | rs298085 |
| rs966220 | rs1545069 | rs1435075 | rs1501640 | rs1504983 | rs298084 |
| rs966221 | rs1545070 | rs1435074 | rs600611 | rs1504982 | rs298083 |
| rs719702 | rs973700 | rs978455 | rs159621 | rs877745 | rs298073 |
| rs2113073 | rs1583434 | rs1827340 | rs159625 | rs877744 | rs298072 |
| rs2113074 | rs1347401 | rs1393083 | rs1435072 | rs2164661 | rs298071 |
| rs2113075 | rs1949017 | rs988364 | rs173945 | rs981230 | rs1421400 |
| rs1035512 | rs723962 | rs1017408 | rs256356 | rs1437124 | rs402874 |
| rs1559277 | rs1355099 | rs2053155 | rs185351 | rs746477 | rs434368 |
| rs1981848 | rs1396473 | rs181923 | rs256355 | rs893191 | rs371011 |
| rs1544788 | rs1369285 | rs1546364 | rs2067024 | rs1992112 | rs298063 |
| rs1544790 | rs1435071 | rs173942 | rs256354 | rs298102 | rs298062 |
| rs1544791 | rs1435070 | rs159616 | rs173944 | rs298101 | rs298061 |
| rs851284 | rs1435083 | rs159620 | rs256353 | rs2164660 | rs298060 |
| rs1396476 | rs991551 | rs1501641 | rs986400 | rs298100 | rs298057 |
| rs1508860 | rs1154790 | rs159619 | rs1504981 | rs298098 | rs298056 |
| rs1974850 | rs1154789 | rs159614 | rs1120533 | rs298096 | rs1370230 |
| rs2136203 | rs714291 | rs159613 | rs256351 | rs298095 | rs297975 |
| rs2174994 | rs981760 | rs159612 | rs190458 | rs298094 | rs297974 |
| rs1508863 | rs1369288 | rs159611 | rs256352 | rs298093 | rs379578 |
| rs1508859 | rs977418 | rs194368 | rs171745 | rs1362942 | rs920190 |
| rs1508864 | rs977417 | rs661576 | rs1157709 | rs1362941 | rs1865962 |
| rs1396474 | rs977416 | rs299627 | rs1910790 | rs298091 | rs298018 |
| rs1543951 | rs1529843 | rs159608 | rs1910789 | rs298090 | rs298021 |
| rs2016324 | rs1529842 | rs159609 | rs1504985 | rs298089 | rs298022 |

| | | | | | |
|----------|-----------|-----------|-----------|-----------|-----------|
| rs298023 | rs2053229 | rs296406 | rs697076 | rs37575 | rs1824154 |
| rs298024 | rs295974 | rs296405 | rs294478 | rs37576 | rs2112911 |
| rs298025 | rs295973 | rs295948 | rs953302 | rs1876209 | rs1551564 |
| rs298026 | rs295972 | rs295947 | rs294479 | rs190486 | rs2034895 |
| rs298027 | rs295971 | rs295946 | rs697075 | rs447261 | rs2081092 |
| rs298028 | rs295970 | rs295945 | rs294481 | rs1506558 | rs2112910 |
| rs298029 | rs295969 | rs295944 | rs294482 | rs1108916 | rs918583 |
| rs298030 | rs295968 | rs1395334 | rs294483 | rs921942 | rs1840838 |
| rs169868 | rs295966 | rs295943 | rs702545 | rs924998 | rs1350298 |
| rs177077 | rs726652 | rs1035321 | rs294484 | rs176705 | rs1990985 |
| rs298032 | rs295965 | rs294494 | rs294485 | rs1156029 | rs1379297 |
| rs298033 | rs1307218 | rs722923 | rs294486 | rs1156028 | rs1817248 |
| rs298034 | rs1307217 | rs294495 | rs702544 | rs931857 | rs244569 |
| rs298035 | rs893190 | rs294496 | rs702543 | rs931856 | rs244568 |
| rs298042 | rs1111495 | rs294497 | rs159194 | rs931855 | rs244567 |
| rs298044 | rs295961 | rs294498 | rs40215 | rs1506557 | rs244565 |
| rs298045 | rs295960 | rs294499 | rs291118 | rs462930 | rs185417 |
| rs298046 | rs295959 | rs294500 | rs1506560 | rs458953 | rs258128 |
| rs298048 | rs295958 | rs294501 | rs37569 | rs174039 | rs258127 |
| rs298049 | rs296410 | rs294503 | rs291119 | rs2174624 | rs258125 |
| rs298050 | rs295957 | rs295936 | rs37571 | rs2135480 | rs1348710 |
| rs298051 | rs295956 | rs1395336 | rs1870077 | rs992726 | rs1348709 |
| rs298052 | rs295955 | rs1395337 | rs159195 | rs294474 | rs1971061 |
| rs298053 | rs295954 | rs294492 | rs37572 | rs294475 | rs1541673 |
| rs190936 | rs295949 | rs159196 | rs37573 | rs988827 | rs1541672 |
| rs298017 | rs295980 | rs159197 | rs167161 | rs988828 | rs258112 |
| rs298016 | rs295979 | rs172362 | rs37574 | rs1350297 | rs258111 |
| rs298015 | rs295978 | rs37579 | rs1506562 | rs1457110 | rs171800 |
| rs298014 | rs1154587 | rs721784 | rs291122 | rs1457111 | rs187716 |

| | | | | | |
|-----------|-----------|-----------|-----------|-----------|-----------|
| rs258110 | rs424839 | rs1118965 | rs35266 | rs255652 | rs26709 |
| rs258109 | rs370891 | rs154028 | rs39672 | rs255650 | rs26710 |
| rs258108 | rs434183 | rs151802 | rs958851 | rs255649 | rs28055 |
| rs258107 | rs444552 | rs244580 | rs244576 | rs2194210 | rs26711 |
| rs665836 | rs433565 | rs1457145 | rs244575 | rs255648 | rs27723 |
| rs392901 | rs1445918 | rs244579 | rs244573 | rs255647 | rs27185 |
| rs383444 | rs441817 | rs255812 | rs35258 | rs154221 | rs27695 |
| rs662643 | rs433161 | rs154029 | rs35259 | rs256752 | rs1445954 |
| rs670169 | rs428059 | rs185333 | rs40121 | rs256120 | rs27549 |
| rs525099 | rs434422 | rs35289 | rs35261 | rs255635 | rs455969 |
| rs669240 | rs427433 | rs35288 | rs35264 | rs185325 | rs26712 |
| rs381755 | rs391377 | rs35287 | rs40122 | rs26686 | rs1867711 |
| rs454702 | rs414746 | rs35286 | rs35265 | rs1031197 | rs1867712 |
| rs443191 | rs187368 | rs35285 | rs35255 | rs1031198 | rs26713 |
| rs380118 | rs244593 | rs35284 | rs721826 | rs27183 | rs26714 |
| rs2168649 | rs244592 | rs35283 | rs244570 | rs28044 | rs27547 |
| rs371775 | rs244591 | rs35282 | rs27171 | rs27182 | rs26715 |
| rs378970 | rs244590 | rs35281 | rs1824159 | rs545611 | rs27949 |
| rs401013 | rs181736 | rs35280 | rs27170 | rs649476 | rs26700 |
| rs427748 | rs193447 | rs35279 | rs27169 | rs1664896 | rs1306348 |
| rs427740 | rs2028842 | rs35278 | rs27168 | rs149106 | rs35309 |
| rs378869 | rs2028841 | rs40126 | rs2013979 | rs1374028 | rs27691 |
| rs1902609 | rs1823068 | rs35277 | rs889231 | rs531105 | rs35310 |
| rs389324 | rs1823067 | rs35276 | rs2014012 | rs27184 | rs26689 |
| rs387647 | rs1823066 | rs35275 | rs37353 | rs1445951 | rs27187 |
| rs377451 | rs244588 | rs40125 | rs187645 | rs1947090 | rs1445948 |
| rs403695 | rs168641 | rs35274 | rs1809012 | rs26708 | rs26687 |
| rs403672 | rs2059175 | rs244577 | rs187644 | rs2112959 | rs166260 |
| rs372309 | rs2059174 | rs35267 | rs153981 | rs1445953 | rs149506 |

| | | | |
|-----------|-----------|-----------|-----------|
| rs27722 | rs1664886 | rs1559251 | rs1553113 |
| rs26695 | rs1867724 | rs1345791 | rs1353748 |
| rs27773 | rs1445947 | rs1345792 | rs1498606 |
| rs1471429 | rs42470 | rs1345793 | rs1353747 |
| rs1471430 | rs1423308 | rs1105577 | rs1006431 |
| rs26705 | rs27174 | rs1960 | rs1948651 |
| rs28054 | rs168834 | rs1824788 | rs1498605 |
| rs26703 | rs27727 | rs1862563 | rs1498604 |
| rs27898 | rs27172 | rs1551939 | rs1498603 |
| rs722010 | rs676449 | rs1038080 | rs1995166 |
| rs27957 | rs27186 | rs997421 | rs1498602 |
| rs26702 | rs2112957 | rs1014317 | rs1077183 |
| rs27548 | rs1023814 | rs2059191 | rs1078368 |
| rs26701 | rs27175 | rs1551938 | rs1874857 |
| rs27188 | rs1445950 | rs1186170 | rs1874858 |
| rs27189 | rs2021384 | rs986067 | rs1909294 |
| rs149084 | rs736736 | rs954740 | rs1546221 |
| rs153968 | rs745813 | rs1363882 | rs2055295 |
| rs464787 | rs889229 | rs1353749 | rs1391648 |
| rs153978 | rs1077978 | rs1391651 | rs2055298 |
| rs464311 | rs2081106 | rs1391650 | rs1472456 |
| rs149108 | rs1559252 | rs1391649 | rs1553114 |
| rs153980 | rs2054443 | rs1391652 | rs1542842 |
| rs153961 | rs922437 | rs950446 | rs1498611 |
| rs1867725 | rs922436 | rs950447 | rs1532520 |
| rs153965 | rs922435 | rs1498599 | |
| rs153966 | rs922434 | rs1498601 | |
| rs1988803 | rs716908 | rs1498609 | |
| rs467300 | rs1971940 | rs1498608 | |

Table 12

New SNPs identified by deCODE

| Position | Variation | AA Change | Exon |
|----------|-----------|-----------|-------------|
| 135641 | T/A | | 1268187 C/T |
| 142780 | A/G | | 1268553 A/G |
| 732790 | G/T | | 1272669 G/A |
| 735966 | C/A | | 1272910 A/G |
| 736226 | A/G | | 1273023 G/A |
| 736516 | C/T | | 1273220 A/G |
| 850001 | G/A | | 1273240 A/G |
| 852776 | A/C | | 1273543 C/T |
| 853079 | G/T | | 1288439 G/A |
| 853575 | C/A | | 1289730 T/A |
| 856468 | A/G | | 1290176 G/A |
| 860845 | A/G | | 1293745 T/C |
| 870924 | A/G | | 1344605 A/G |
| 1027267 | T/C | | 1344864 G/A |
| 1027643 | T/G | | 1345135 C/G |
| 1027757 | T/C | | 1345286 A/G |
| 1028146 | T/A | | 1346112 C/T |
| 1037657 | A/C | | 1352976 A/T |
| 1044016 | G/A | | 1354291 T/C |
| 1044045 | C/T | | 1354377 C/T |
| 1254737 | T/C | | 1354554 C/A |
| 1254849 | T/C | | 1354675 T/C |
| 1255763 | G/T | | 1355114 T/C |
| 1257206 | A/G | | 1355693 A/G |
| 1258161 | T/C | | 1357081 A/G |
| 1268007 | A/G | | 1362985 T/G |

| | | | | | |
|---------|-----|---------|-----|----------|--------|
| 1363021 | C/T | 1580088 | G/A | | |
| 1363827 | C/T | 1581078 | G/A | | |
| 1363911 | G/A | 1582418 | T/A | | |
| 1364061 | C/T | 1584580 | A/C | | |
| 1364066 | T/A | 1585955 | G/T | | |
| 1367904 | A/G | 1590608 | T/C | | |
| 1368193 | T/C | 1590672 | A/G | | |
| 1368217 | G/C | 1590673 | G/T | | |
| 1373349 | C/T | 1590837 | G/A | | |
| 1373384 | A/G | 1590936 | C/A | | |
| 1373415 | T/C | 1591011 | G/A | | |
| 1373979 | T/G | 1591047 | C/T | | |
| 1376149 | G/A | 1591306 | C/A | Pro->Thr | D1 |
| 1384931 | A/C | 1591583 | T/C | | |
| 1385093 | A/T | 1594788 | C/A | | |
| 1385107 | G/A | 1594994 | G/A | | |
| 1385445 | T/C | 1601831 | C/T | | |
| 1391418 | G/C | 1636902 | T/C | | |
| 1409210 | C/A | 1638550 | A/C | Lys->Thr | exon 4 |
| 1414804 | C/T | 1640663 | T/C | | |
| 1428284 | T/C | 1641954 | C/T | | |
| 1431800 | A/T | 1641960 | C/T | | |
| 1449904 | A/T | 1653881 | G/A | | |
| 1574301 | C/G | 1655748 | G/A | | |
| 1574615 | C/T | 91470 | G/A | | |
| 1575634 | A/T | | | | |

Discussion of Example 2:

5 Here we present the first complete genomic sequence of human PDE4D, two
novel mRNA/protein isoforms of PDE4D and their corresponding exons, and the intron-
exon structure of known and novel isoforms. The basis for phosphodiesterases is the
mammalian homolog of the "dunce" gene in *Drosophila melanogaster*, implicated in
learning and memory (Davis, R.L. and B. Dauwalder, *Trends Genet.*, 7(7):224-229
10 (1991)). PDEs are members of a large superfamily of isoenzymes subdivided into 9 and
possibly 10 distinct families (Conti, M. and S.L. Jin, *Prog. Nucleic Acid Res. Mol. Biol.*,
63:1-38 (1999)), with several genes in each family and more than one isoform for each
gene. The significance of the diversity of PDEs is not known but many of the isoforms
differ in their biochemical properties, phosphorylation, intracellular targeting, protein-
15 protein interactions and patterns of expression in tissues, which suggests that each of the
various isoforms might have distinct functions (Bolger, G.B., *Cell Signal*, 6(8):851-859
(1994); Conti, M., *et al.*, *Endocr. Rev.*, 16(3):370-378 (1995)).

There are four genes that encode the type 5 PDEs (PDE4A, PDE4B, PDE4C and
PDE4D), which is a group of enzymes characterized by high affinity for cAMP. The
20 gene for PDE4D was assigned to human chromosome 5q12 (Milatovich, A., *et al.*,
Somat. Cell Mol. Genet., 20(2):75-86 (1994); Szpirer, C., *et al.*, *Cytogenet. Cell Genet.*,
69(1-2):22-14 (1995)) and 5 distinct splice variants have been characterized (the short
forms PDE4D1, PDE4D2 and the long forms PDE4D3, PDE4D4, and PDE4D5)
(Bolger, G.B., *et al.*, *Biochem. J.*, 328(Pt.2):539-548 (1997)) (FIG. 3). The sequence of
25 the human PDE4D variants show a high degree of homology to the PDE4Ds expressed
in mouse and rat. The pattern of splicing and different promoter usage is highly
conserved during evolution indicating an important physiological role (Nemoz, G., *et*
al., *FEBS Lett.*, 384(1):97-102 (1996)). The PDE4D variants are generated at two major
boundaries present in the gene. The first boundary corresponds to the junction of exon
30 2. Differential splicing in this region generates the 2 short variants PDE4D1 (586 a.a.)

and PDE4D2 (508 a.a.) (FIG. 3). This splicing boundary is conserved in mouse, rat and between different human PDE4 genes. The splicing variant PDE4D2 is generated by the removal of 256 bp from the PDE4D1 sequence. The initiation codon in the PDE4D2 variant lies within exon D1/D2. Data demonstrates that the expression of the short
5 PDE4D variants is under the control of an internal promoter regulated by cAMP (Vicini, E. and M. Conti, *Mol. Endocrinol.*, 11(7):839-850 (1997)). The second major splicing boundary is also conserved during evolution and is identical to that described in the *Drosophila dunce* gene. Splicing occurs at the intron/exon boundary at the LF1 exon (FIG. 3).

10

PDE function

The PDEs serve at least four major functions in the cell. They can (1) act as effector of signal transduction by interacting with receptors and G-proteins; (2) integrate the cyclic nucleotide-dependent pathway with other signal transduction pathways; (3)
15 function as homeostatic regulators, playing a role in feedback mechanisms controlling cyclic nucleotide levels during hormone and neurotransmitter stimulation; (4) play an important role in controlling the diffusion of cyclic nucleotides and in creating subcellular domains or channeling cyclic nucleotide signaling (Conti, M. and S.L. Jin, *Prog. Nucleic Acid Res. Mol Biol.*, 63:1-38.(1999)). Inhibition of PDE has long been
20 recognized as an effective pharmacological strategy to alter intracellular cyclic nucleotide levels (Flamm, E.S., *et al.*, *Arch. Neurol.*, 32(8):569-71 (1975)).

It has been reported that PDE4 is the predominant isozyme regulating vascular tone mediated by cAMP hydrolysis in cerebral vessels (Willette, R.N., *et al.*, *J. Cereb. Blood Flow Metab.*, 17(2):210-9 (1997)).

25 A recent study on mice with targeted disruption of PDE4D gene (Hansen, G., *et al.*, *Proc. Natl. Acad. Sci. U S A*, 97(12):6751-6 (2000)) has demonstrated a crucial role of PDE4D in the control of smooth muscle contraction and muscarinic cholinergic receptor signaling but not in the control of airway inflammation. The lung phenotype of the PDE4D^{-/-} mice demonstrates that this gene plays a nonredundant role in cAMP

homeostasis. There is a significant reduction in PDE activity and an increase in resting and stimulated cAMP levels in the lung, indicating that other PDE4s (or other PDEs) are not up-regulated and cannot compensate for the loss of PDE4D. These findings support that PDE4D serves a unique, nonoverlapping functions in cell signalling.

- 5 No clear link between an established inherited disorder and known PDE loci has emerged, with the exception of PDE6. Inhibitors of PDEs have been shown to affect airway responsiveness and pulmonary allergic inflammation (Schudt, C., *et al.*, *Pulm. Pharmacol. Ther.*, 12(2):123-9 (1999)). There are reports suggesting that altered PDE4 function may be linked to nephrogenic diabetes insipidus (Takeda, S., *et al.*,
10 *Endocrinology*, 129(1):287-94 (1991)) or atopic dermatitis (Chan, S.C., *et al.*, *J. Allergy Clin. Immunol.*, 91(6):1179-88 (1993)), however no mutations have been identified. It has also been reported that vasorelaxation modulated by PDE4 (not mentioned whether it is A, B, C or D gene family) is compromised in chronic cerebral vasospasm associated with subarachnoid hemorrhage (Willette, R.N., *et al.*, *J. Cereb. Blood Flow Metab.*,
15 17(2):210-9 (1997)). PDE4D itself has not been linked to stroke before.

PDE4D expression and cellular localization

- PDE4Ds are expressed in human peripheral mononuclear cells (Nemoz, G., *et al.*, *FEBS Lett*, 384(1):97-102 (1996)), brain (Bolger, G., *et al.*, *Mol. Cell Biol.*,
20 13(10):6558-71 (1993)), heart (Kostic, M.M., *et al.*, *J. Mol. Cell Cardiol.*, 29(11):3135-46 (1997)) and vascular smooth muscle cells (Liu, H. and D.H. Maurice, *J. Biol. Chem.*, 274(15):10557-65 (1999)).

- Immunoblotting of rat brain has shown that the PDE4D3, PDE4D4 and PDE4D5 proteins are present in brain (Bolger, G.B., *et al.*, *Biochem. J.*, 328(Pt 2):539-48 (1997))
25 and are expressed in cortex and cerebellum from rat (Iona, S., *et al.*, *Mol. Pharmacol.*, 53(1):23-32 (1998)). These proteins were recovered mostly or exclusively in the particulate fraction suggesting that these forms may be targeted to insoluble cellular structures. In addition a 68 kDa protein was detected which could represent PDE4D1, PDE4D2 or both. To verify this RT-PCR was performed on mRNA from rat brain and

the results showed that transcripts for PDE4D1 and 2 were present. Their data also suggests that the N-terminal regions of the PDE4D3-5, derived from alternatively spliced regions of their mRNAs, are important in determining their subcellular localization activity and differential sensitivity to inhibitors and there are indications that there is a propensity for the long PDE4D isoforms to interact with particulate fraction of the cell.

EXAMPLE 3: PDE4D ISOFORM EXPRESSION

10 *Expression analysis in EBV transformed B cell lines*

As a functional mutation in the known coding exons of *PDE4D* was not identified, gene expression was next studied to determine if the genetic association to stroke relates to regulation of its expression levels. In order to test this, we chose to use cell lines instead of blood or tissues for these studies because expression analysis of cell lines is not confounded by the presence of multiple cell types. Cell types may express *PDE4D* at different levels so it is generally more reliable to quantify expression in cell lines than tissues. Isoform-specific kinetic PCR analysis was carried out on EBV transformed B cell lines to quantify each isoform in 83 stroke patients and 84 controls. These patients were not selected for this analysis based on any specific subtype of stroke. The majority of the patients had ischemic stroke and 38% of them had carotid or cardiogenic cause of stroke. Overall the total PDE4D message level as assessed by amplification across exons present in all isoforms (PAN), was significantly lower in patients than in controls (p value < 0.005). This decrease was due primarily to lower expression of the isoforms, PDE4D1, PDE4D2 and PDE4D5 (FIG. 4).

25 We selected individuals with a specific stroke associated haplotype and compared the expression levels of carrier vs. non-carriers of this haplotype and with patients and controls examined separately (FIGS. 5 and 6). The haplotype was constructed out of the at-risk allele for the microsatellite marker AC008818-1 and SNP45 (SNP5PDM357221) and SNP41 (SNP5PDM361545). This haplotype acts as a

surrogate for the disease-associated haplotype we have identified in LD block B (Table 3). Patients with the haplotype had a significantly decreased expression of the PDE4D7 and PDE4D9 isoforms (FIG. 5). Several other isoforms of PDE4D were expressed but did not show correlation to the disease haplotype. The PDE4D7 correlation was also present in controls but only marginally significant (FIG. 6). Of interest, this at-risk haplotype covers the 5' exon specific to PDE4D7 and presumably its promoter.

These results show that there is significant dysregulation of the expression of multiple PDE4D isoforms in stroke patients.

10 *Methodology for expression analysis using Quantitative Reverse Transcriptase PCR*

Total RNA was isolated from EBV transformed B-cell cultures according to manual, using the TRIZOL[®] reagent provided by GibcoBRL. RNeasy mini Qiagen kit with on column DNA digestion was used to clean RNA. Quality and quantity of RNA was assessed using 2100 Agilent Bioanalyser. cDNA was prepared from total RNA using random hexamers with TaqMan Reverse Transcription Reagents kit from Applied Biosystems (N808-0234). Primer Express 2.0 and Oligo 6 software were used to make cDNA specific primers and probes for PDE4D and PDE4D isoforms. GAPDH "Assay-On-Demand" was obtained from Applied Biosystems and used as a housekeeping gene. PDE assays were tested and optimized for 384 well high throughput expression analysis using ABI 7900 Instrument. A final concentration of 200 nM probes, 900 nM primers and 2 ng/mcl cDNA was used in a 10mcl reaction volume. Each plate was run twice and an average for each sample calculated. ABI7900 instrument was used to calculate CT (Threshold Cycle) values. Samples displaying a greater than 1 deltaCT between duplicates were not used in our analysis. Quantity was obtained using the formula $2^{-\Delta CT}$ where ΔCT represents the difference of CT values between target and housekeeping assay.

Accession numbers AY245866 (PDE4D7) and AY245867 (PDE4D9).

Discussion of the three examples and conclusions:

Our results indicate that genetic variation in the *PDE4D* gene is associated with ischemic stroke. The direct involvement of *PDE4D* is strongly supported by linkage in conjunction with association and expression analysis. We first identified the association
5 using microsatellite markers, and supplementing the microsatellite data with a denser set of SNPs further supported this. The strongest association is to the two ischemic subtypes, carotid and cardiogenic stroke whereas we did not observe association to small vessel occlusive disease, the form of stroke thought to be independent of atherosclerosis. Although we have not identified a functional mutation in the *PDE4D* gene, we have
10 identified a haplotype, that extends over the first exon of *PDE4D* that is significantly associated to carotid and cardiogenic stroke. This haplotype is present in 47% of the carotid/cardiogenic stroke patients, compared to 21% in the control group with more than two-fold stroke risk for the carriers of this haplotype. It has a population attributed risk of 25%. For the combined cardiogenic and carotid subtype of stroke, apart from
15 finding individual SNP and microsatellite alleles that are significantly associated with the disease even after adjusting for multiple comparison, most interesting is the discovery that haplotypes covering the first exon of *PDE4D* can be classified into three groups with clearly distinct risks. Relative to the protective group, the population attributed risk of the at-risk and wild type groups combined is estimated to be 55%.
20 Approximately 16% of the population carries one copy of the at-risk haplotype in FIG. 12.3. They have about 1.8 times the risk of the general population for getting cardiogenic or carotid stroke. Approximately 0.8% of the population is homozygous for the at-risk haplotype and, assuming the multiplicative model, their risk is estimated to be about 3.8 times the risk of the general population. It is true that we have not yet
25 identified or proved convincingly what is the functional variant, or variants, which are responsible for the observed effects of these haplotype groups. And, since these haplotype groups do not fully explain the linkage signal we observe in the region for all stroke patients, we certainly could not rule out, and indeed expect, that there are other variants/haplotypes within *PDE4D* not directly related to those we have identified that

confer risk to stroke. These are likely to be rare but could have very high penetrance. We also cannot rule out the possibility that some other genes in the linkage region independent of, or in conjunction with, *PDE4D* confer susceptibility to stroke.

We examined whether the disease associated alleles and haplotype are related to specific stroke risk factors such as hypertension, hypercholesterolemia, diabetes, peripheral artery occlusive disease and coronary artery disease in addition to each onset of stroke and gender (Table 8). A marginally significant association to hypercholesterolemia was observed but it is clear that *PDE4D*'s contribution to stroke is not strongly correlated with any of these known risk factors.

The *PDE4D* gene is a highly complex gene. By alternative splicing and use of different promoters this gene generates at least 8 different isoforms that yield functional proteins, differing from each other in their N-terminal regions. We have identified four new exons encoding the N-termini of two new isoforms *PDE4D7* and *PDE4D9*. The disease-associated haplotype extends over the 5'exon unique to the new *PDE4D7* variant and the presumed promoter region of this isoform suggesting that the functional variation may be involved in transcriptional regulation. This hypothesis is also supported by our *PDE4D* expression analysis that shows significant correlation between the disease associated haplotype and the level of *PDE4D7* message.

The strongest association found for this *PDE4D* haplotype was to the two major subtypes of ischemic stroke, carotid and cardiogenic stroke, suggesting a role for this gene in the vascular biology of atherosclerosis. While there are multiple etiologies for ischemic stroke, atherosclerosis remains the most important one and it is the major pathological process for the two ischemic subtypes, carotid and cardiogenic strokes. First, it is the major cause of stenotic and occlusive lesions of the internal and common carotids that lead to carotid strokes. Second, cardiac thrombi which shed emboli to the brain most commonly occur on the background of coronary artery disease, such as following acute myocardial infarction or ischemic cardiomyopathy, and/or due to atrial fibrillation on the basis of poor compliance of ischemic ventricles (diastolic dysfunction/stiffening). Although atrial fibrillation may occur on the background of

other diseases such as valvular disease, hyperthyroidism, and hypertension, in the age group that tends to suffer from stroke, ischemic heart disease remains one of the most important causes. Ischemic stroke resulting from occlusion of small penetrating arteries within the brain (small vessel occlusive disease or lacunar stroke) is generally thought to result from endothelial proliferation since atherosclerosis only occurs in larger arteries. PDE4D does not show association to small vessel stroke, consistent with its role in atherosclerosis. Carotid and cardiogenic stroke together account for the majority of ischemic stroke (note that our number for carotid is lower since we used a more stringent cutoff of stenosis).

10 PDE4D selectively degrades second messenger cAMP (Kong, A. *et al.*, *Nat Genet* 10, 10 (2002)), which plays a central role in signal transduction and regulation of physiological responses. It is expressed in most cell types important to the pathogenesis of atherosclerosis, including vascular smooth muscle cells (VSCM), endothelial cells, monocytes, macrophages and T-lymphocytes (Houslay, M.D. and Adams, D.R.,
15 *Biochem J* 370, 1-18 (2003); Liu, H. and Maurice, D.H., *J Biol Chem* 274, 10557-65. (1999); Liu, H. *et al.*, *J Biol Chem* 275, 26615-24. (2000); Baillie, G., *et al.*, *Mol Pharmacol* 60, 1100-11. (2001); Jin, S.L. and Conti, M., *Proc Natl Acad Sci US A* 99, 7628-33. (2002)). Cyclic AMP is a key signalling-molecule in these cells (Landells, L.J. *et al.*, *Br J Pharmacol* 133, 722-9 (2001); Fukumoto, S. *et al.*, *Circ Res* 85, 985-91. (1999); Ogawa, S. *et al.*, *Am J Physiol* 262, C546-54 (1992)). In VSMC, low cAMP
20 levels lead to an increase in proliferation and migration that at least in part is mediated by PDE4 (Landells, L.J. *et al.*, *Br J Pharmacol* 133, 722-9 (2001); Stelzner, T.J., *et al.*, *J Cell Physiol* 139, 157-66 (1989); Pan, X., *et al.*, *Biochem Pharmacol* 48, 827-35. (1994)). Animal models have also shown that elevation of cAMP reduces neointimal
25 lesion formation and inhibits proliferation of SMCs after arterial injury (Palmer, D., *et al.*, *Circ Res* 82, 852-61. (1998); Indolfi, C. *et al.*, *Nat Med* 3, 775-9. (1997)). In monocytes and T-lymphocytes, accumulation of cAMP is generally associated with inhibition of immune functions such as proliferation and cytokine secretion (Indolfi, C. *et al.*, *J Am Coll Cardiol* 36, 288-93. (2000)). It is attractive to postulate that the

regulation of cAMP through absolute or relative expression of one or more PDE4D isoforms may differ in individuals susceptible to stroke; some stroke patients may have increased PDE4D activity and, consequently lower cAMP levels in any of the above cell types, leading to development of the atherosclerotic plaque and/or its instability.

5 However, contrary to what one might expect we see decreased expression in some of the PDE4D isoforms in EBV cell lines from stroke patients. It is of interest that these isoforms are all up regulated by cAMP (Liu, H. and Maurice, D.H., *J Biol Chem* 274, 10557-65. (1999); Tilley, S.L., *et al.*, *J Clin Invest* 108, 15-23 (2001); Vicini, E. and Conti, M., *Mol Endocrinol* 11, 839-50 (1997)) suggesting dysregulation at the level of
10 cAMP in patients. It is therefore possible that increased activity of one or few splice variants alters the effective PDE4D enzymatic activity of the cell decreasing the cAMP levels thus altering the expression of cAMP regulated isoforms as observed in our expression study. This relative expression of PDE4D isoforms may determine the compartmental localization of PDE4D isoforms and thus the corresponding gradients of
15 intracellular cAMP that have been recently observed (see Housley review).

In summary, we have presented association analyses (single marker and haplotype analyses) that support the notion that the *PDE4D* gene confers risk to ischemic stroke. Furthermore, we have observed significant dysregulation of multiple PDE4D isoforms in stroke patients. We propose that this gene is involved in the
20 pathogenesis of stroke through atherosclerosis. PDE4D is expressed in cell types important in atherosclerosis and regulates a second messenger with a central role to processes important in the pathogenesis of atherosclerosis. Inhibition of PDE4D in general or specifically one or more isoforms, by a small molecule drug or other pharmacological agent might decrease the risk of stroke in general, and especially those
25 who are predisposed to stroke through variation in the *PDE4D* gene.

While this invention has been particularly shown and described with reference to preferred embodiments thereof, it will be understood by those skilled in the art that various changes in form and details may be made therein without departing from the spirit and scope of the invention as defined by the appended claims.

CLAIMS

What is claimed is:

- 5 1. A method of diagnosing susceptibility to a stroke in an individual, comprising screening for an at-risk haplotype in the phosphodiesterase 4D gene that is more frequently present in an individual susceptible to stroke compared to a healthy individual, wherein the at-risk haplotype increases risk of stroke significantly.
- 10 2. The method of claim 1 wherein the significant increase is at least about 20%.
3. The method of claim 1 wherein the significant increase is identified as an odds ratio of at least about 1.2.
- 15 4. A method of diagnosing susceptibility to stroke in an individual, comprising screening for an at-risk haplotype in the phosphodiesterase 4D gene that is more frequently present in an individual susceptible to stroke (affected), compared to the frequency of its presence in a healthy individual (control), wherein the presence of the at-risk haplotype is indicative of a susceptibility to stroke.
- 20 5. The method of Claim 4 wherein the at risk haplotype 1 is characterized by the presence of G at nucleic acid position 142780, relative to SEQ ID NO: 1 and allele 0 of microsatellite marker AC0088181-1.
- 25 6. The method of Claim 4 wherein the at risk haplotype 2 is characterized by the presence of G T A A C C A C G A A C T T A T T G A A T T T G A A at nucleic acid positions: 142780, 135112, 132562, 131865, 129361, 129360, 125304, 123426, 123312, 120628, 118914, 111781, 111252, 109301, 107849, 105225, 104552, 102977, 100795, 99035, 88614, 88456, 83119, 82244, 80127,

78552, relative to SEQ ID NO: 1 and allele 0 of microsatellite marker AC0088181-1.

- 5 7. The method of Claim 4 wherein the at risk haplotype 3 is characterized by the presence of A A C A A at nucleic acid positions 138806, 131865, 129361, 120628, 91470, relative to SEQ ID NO: 1.
- 10 8. The method of Claim 4 wherein screening for the presence of an at-risk haplotype within or near PDE4D that significantly correlates with haplotype 1 or stroke susceptibility.
- 15 9. The method of Claim 4 wherein screening for the presence of an at-risk haplotype within or near PDE4D that significantly correlates with haplotype 2 or stroke susceptibility.
- 20 10. The method of Claim 4 wherein screening for the presence of an at-risk haplotype within or near PDE4D that significantly correlates with haplotype 3 or stroke susceptibility.
- 25 11. The method of Claim 4 wherein screening for the presence of an at-risk haplotype in the phosphodiesterase 4D gene comprises enzymatic amplification of nucleic acid from said individual.
12. The method of Claim 11 wherein the nucleic acid is DNA.
13. The method of Claim 12 wherein the DNA is mammalian.
14. The method of Claim 13 wherein the DNA is human.

15. The method of Claim 4 wherein screening for the presence of an at-risk haplotype in the phosphodiesterase 4D gene comprises:
- (a) obtaining material containing nucleic acid from the individual;
 - (b) amplifying said nucleic acid; and
 - 5 (c) determining the presence or absence of an at-risk haplotype in said amplified nucleic acid.
16. The method of Claim 15 wherein determining the presence of an at-risk haplotype is performed by electrophoretic analysis.
- 10 17. The method of Claim 15 wherein determining the presence of an at-risk haplotype is performed by restriction length polymorphism analysis.
18. The method of Claim 15 wherein determining the presence of an at-risk haplotype is performed by sequence analysis.
- 15 19. The method of Claim 15 wherein determining the presence of an at-risk haplotype is performed by hybridization analysis.
- 20 20. A kit for diagnosing susceptibility to stroke in an individual comprising: primers for nucleic acid amplification of a region of the phosphodiesterase 4D gene comprising an at-risk haplotype.
21. The kit of Claim 20 wherein the primers comprise a segment of nucleic acids of length suitable for nucleic acid amplification a single nucleotide polymorphism at nucleic acid position 142780 respectively, relative to SEQ ID NO: 1 and allele 0 of microsatellite marker AC0088181-1.
- 25

22. The kit of Claim 20 wherein the primers comprise a segment of nucleic acids of length suitable for nucleic acid amplification, selected from the group consisting of: single nucleotide polymorphism or microsatellite marker at nucleic acid position 142780, 135112, 132562, 131865, 129361, 129360, 125304, 123426, 123312, 120628, 118914, 111781, 111252, 109301, 107849, 105225, 104552, 102977, 100795, 99035, 88614, 88456, 83119, 82244, 80127, 78552, relative to SEQ ID NO: 1, allele 0 of microsatellite marker AC0088181-1. and combinations thereof.
23. The kit of Claim 20 wherein the primers comprise a segment of nucleic acids of length suitable for nucleic acid amplification, selected from the group consisting of: single nucleotide polymorphism at nucleic acid position at nucleic acid position 138806, 131865, 129361, 120628, 91470, relative to SEQ ID NO: 1 and combinations thereof.
24. A method for assessing susceptibility to stroke in an individual, comprising determining PDE4D isoform expression levels in the individual compared to control, wherein a difference in isoform expression is indicative of susceptibility to stroke.
25. The method of Claim 24 wherein isoform PDE4D7 and/or PDE4D9 expression is determined.
26. A method of diagnosing a susceptibility to stroke, comprising detecting an alteration in the expression or composition of a polypeptide encoded by phosphodiesterase 4D gene in a test sample, in comparison with the expression or composition of a polypeptide encoded by phosphodiesterase 4D gene in a control sample, wherein the presence of an alteration in expression or

composition of the polypeptide in the test sample is indicative of a susceptibility to stroke.

27. The method of Claim 26, wherein the alteration in the expression or composition of a polypeptide encoded by phosphodiesterase 4D gene comprises expression of a splicing variant polypeptide in a test sample that differs from a splicing variant polypeptide expressed in a control sample.
28. A method for preventing the occurrence of stroke in an individual in need thereof, comprising regulating a PDE4D isoform level compared to control, whereby the regulated isoform level mimics the level in a healthy individual.
29. The method of Claim 28 wherein isoform level is regulated by regulating expression of the isoform using a phosphodiesterase 4D gene binding agent, a phosphodiesterase 4D gene receptor, a peptidomimetic, a fusion protein, a prodrug, an antibody or a ribozyme.
30. The method of Claim 28 wherein the isoform level is controlled by genetically altering the isoform's expression level.
31. The method of Claim 28 wherein the isoform level is regulated by altering the ratio of isoforms.
32. The method of Claim 28 wherein isoform PDE4D7 and/or PDE4D9 is regulated.
33. A method for monitoring the effectiveness of treatment on the regulation of expression of one or more PDE4D isoforms at the RNA or protein level, or its enzymatic activity by measuring PDE4D message or protein or enzymatic activity in a sample of peripheral blood or cells derived thereof.

34. A method for predicting the effectiveness of a given therapeutic for stroke prevention or treatment in a given individual comprising screening for the presence or absence of the stroke at-risk haplotype in the phosphodiesterase 4D gene.
35. A method for predicting the effectiveness of a given therapeutic for stroke prevention or treatment in a given individual comprising screening for the expression of one or more PDE4D isoforms at the RNA or protein level, or its enzymatic activity by measuring PDE4D message or protein or enzymatic activity in a sample of peripheral blood or cells derived thereof.
36. A method of diagnosing a reduced or protective susceptibility to a stroke in an individual, comprising screening for a protective haplotype in the phosphodiesterase, 4D gene that is more frequently present in an individual compared to an individual susceptible to stroke, wherein the protective haplotype decreases the risk of stroke significantly
37. A method of Claim 36 wherein the protective haplotype is characterized by the A allele at position 142780, relative to SEQ ID NO: 1 and allele -8 for microsatellitemarker AC0088181-1.

Fig. 1.1

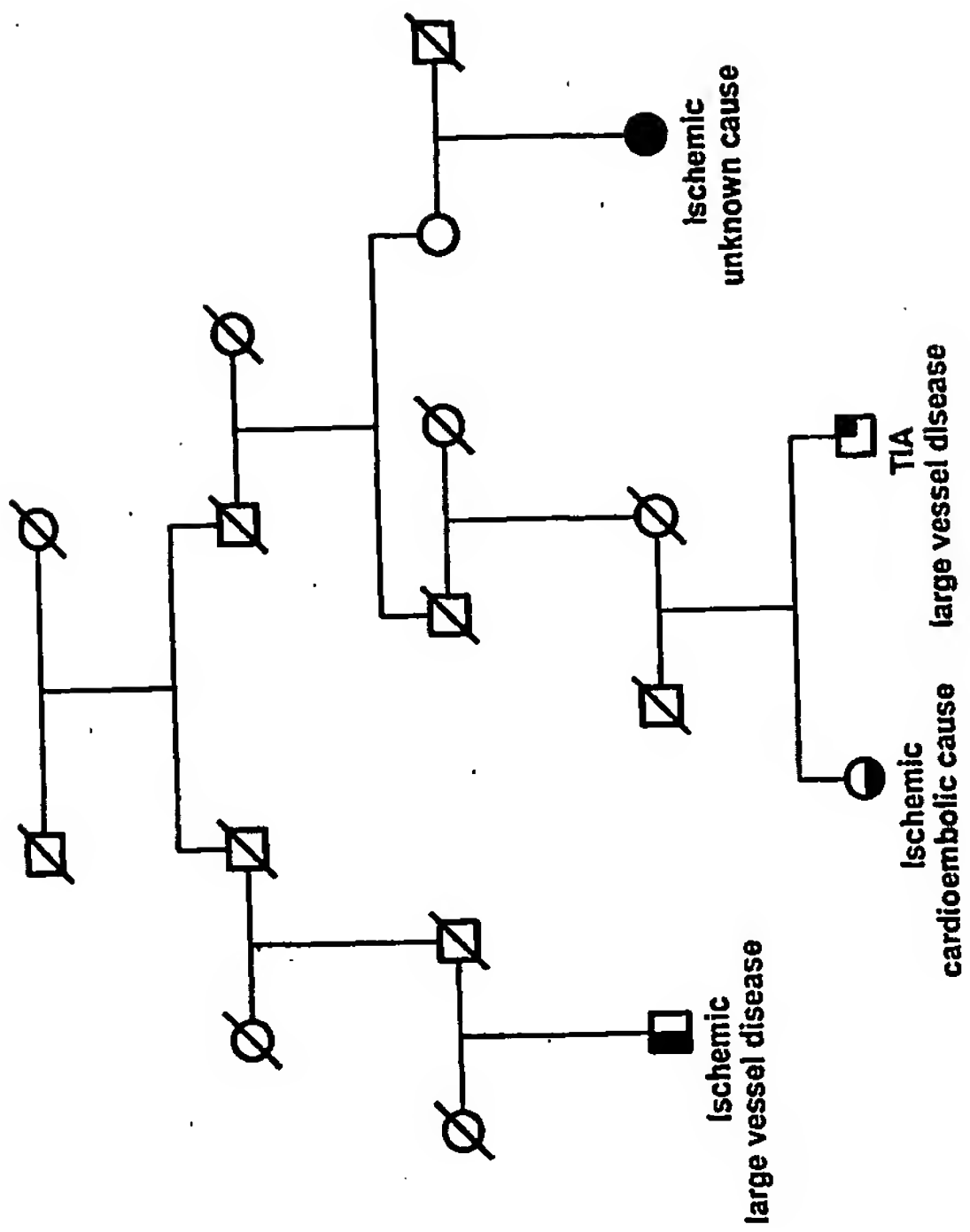
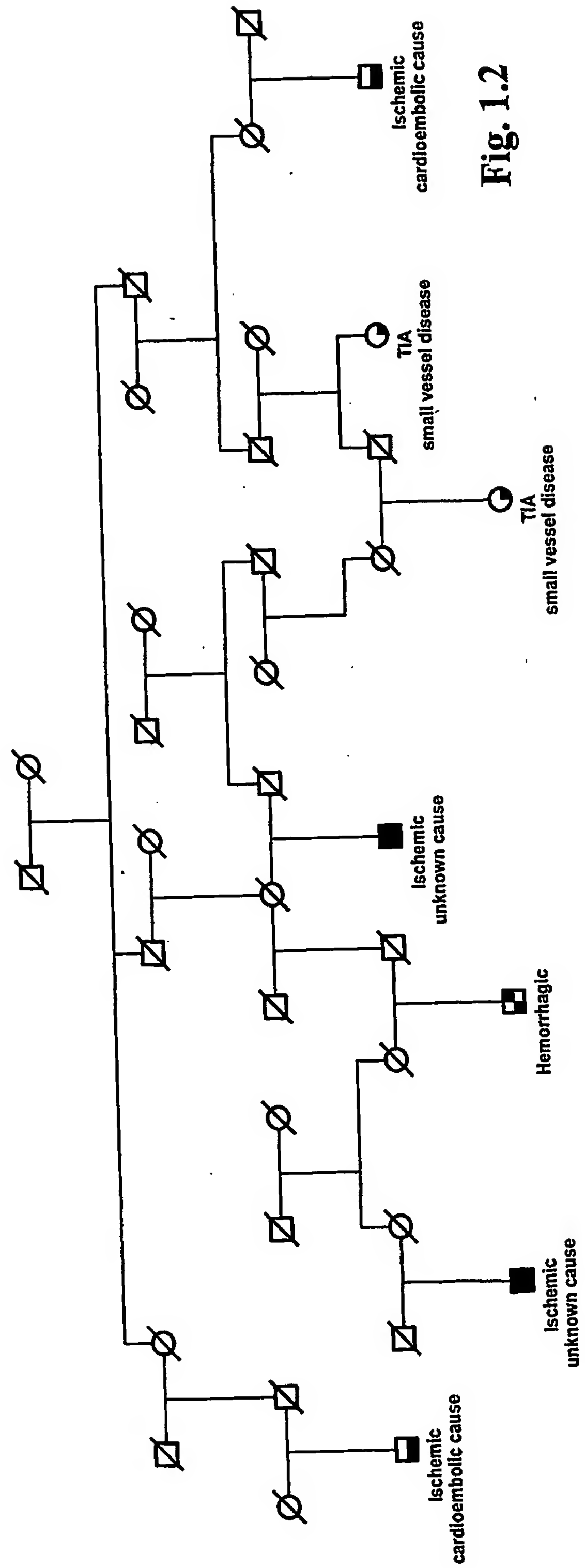


Fig. 1.2



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Genetic map

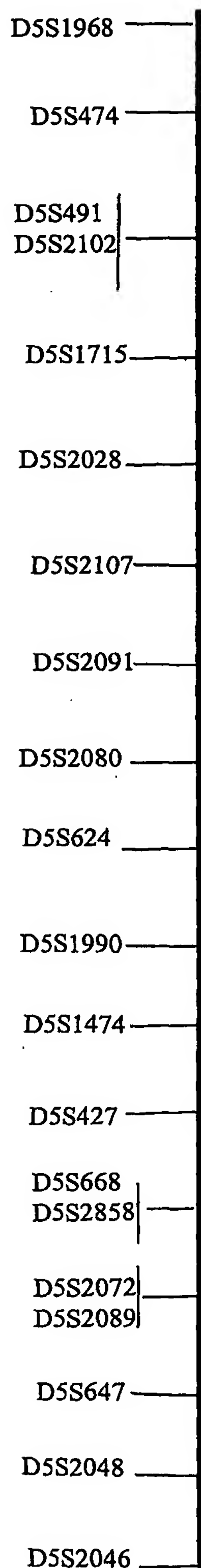
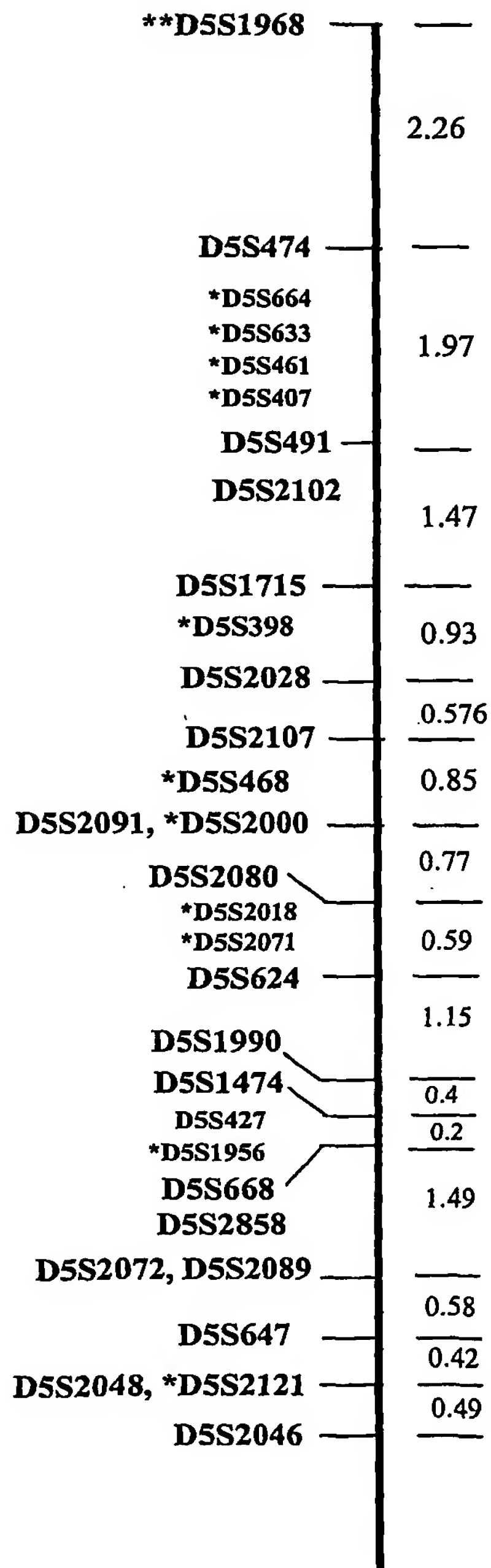


Fig. 2.1

Combined map - cM



*Markers only assigned in physical map

**Marker in blue - only assigned in genetic map

Fig. 2.2

Physical map - Mb

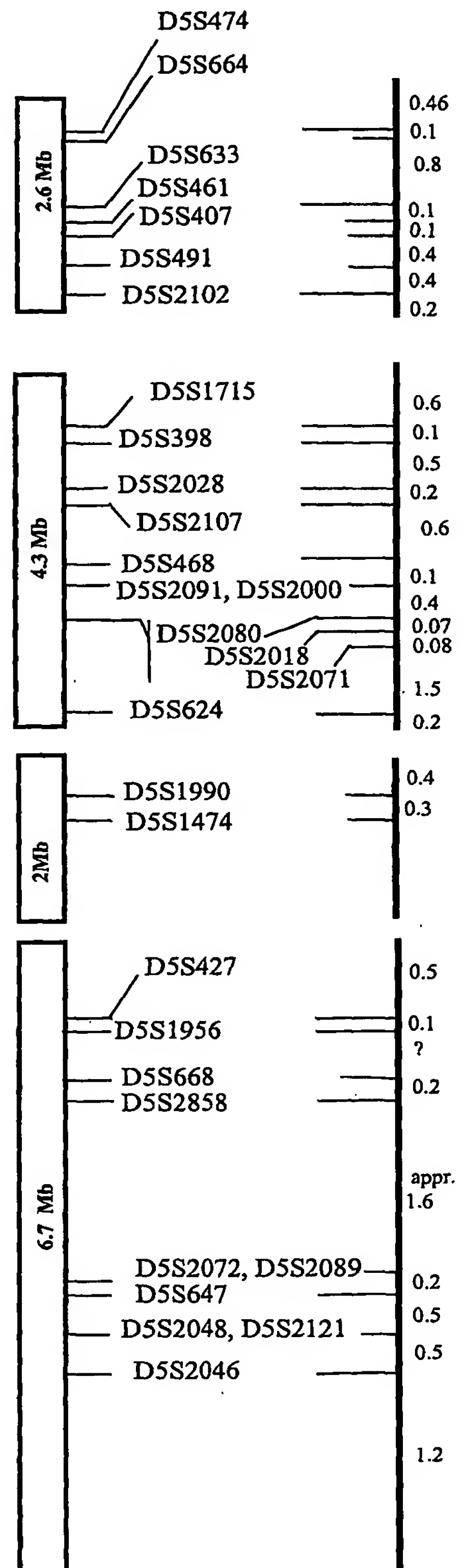


Fig. 2.3

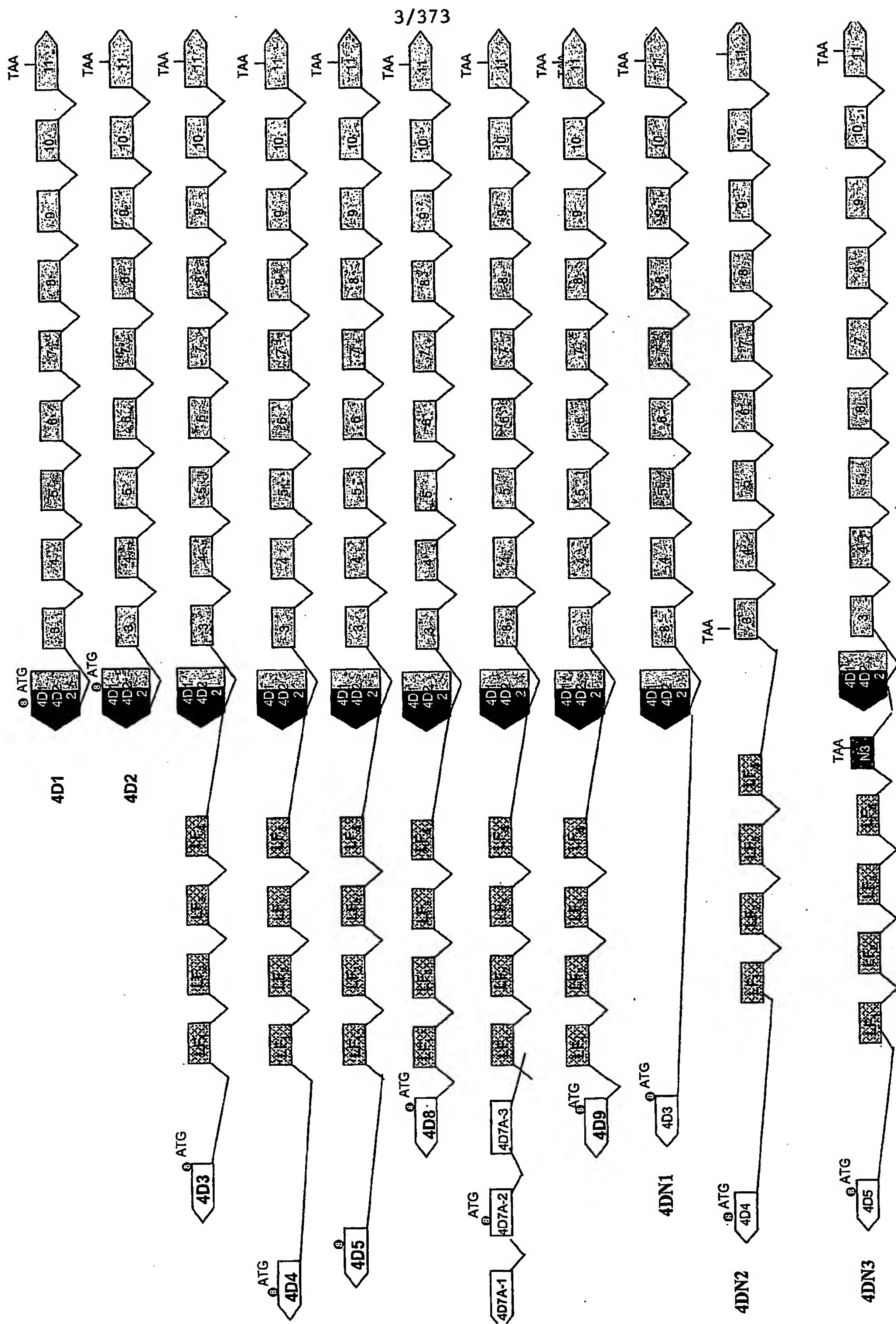
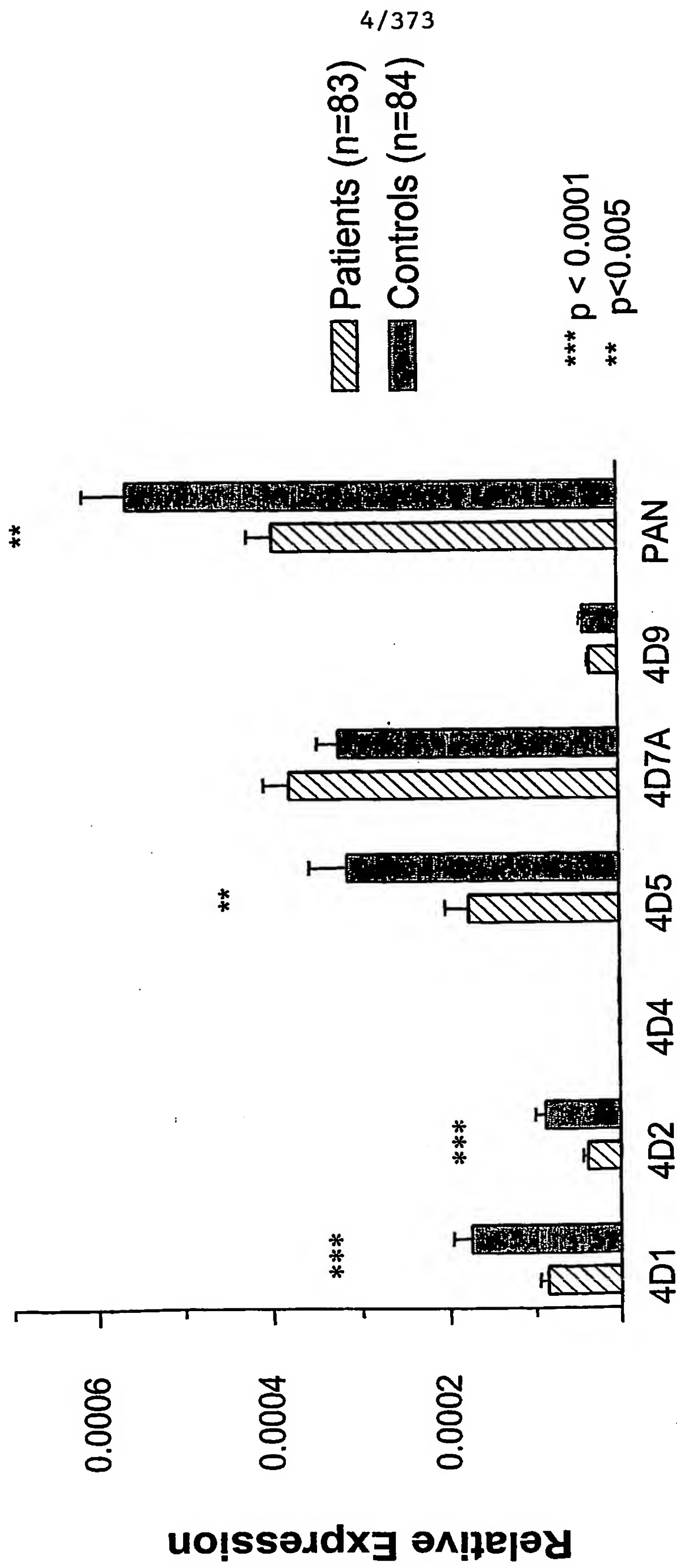


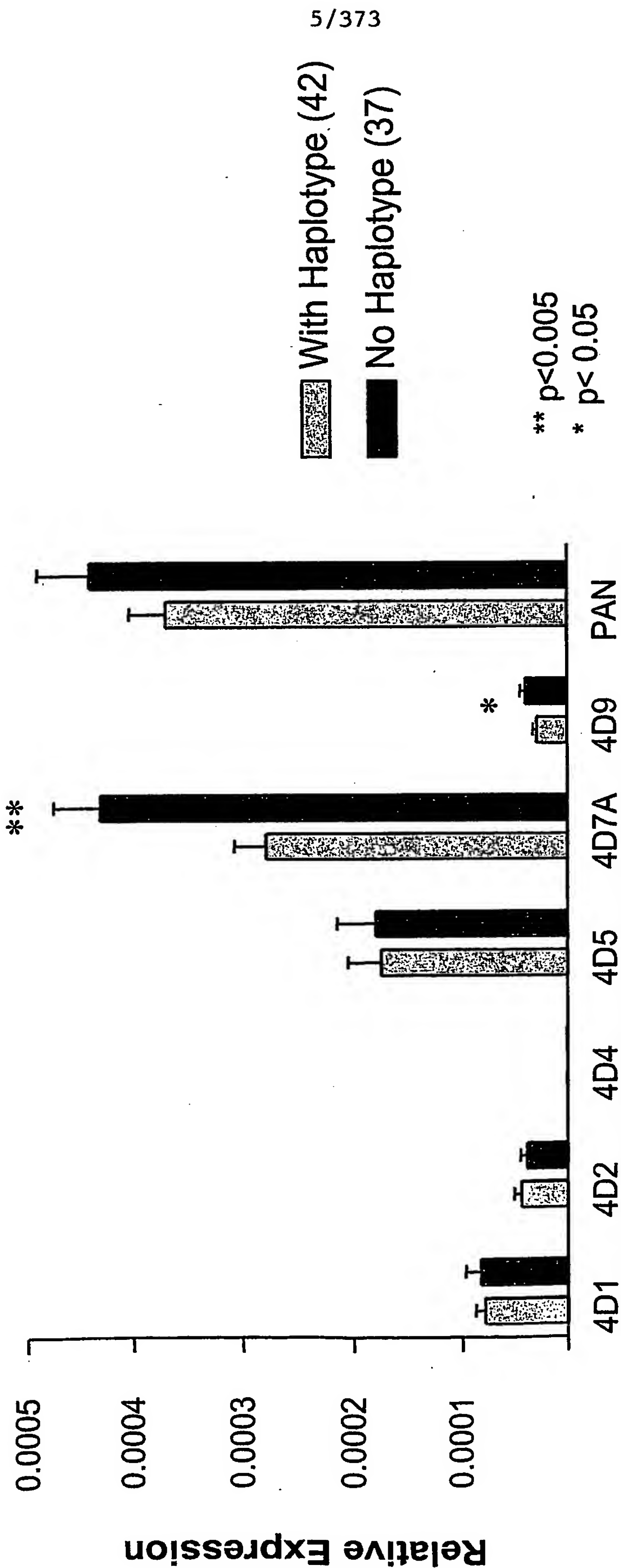
FIG. 3



PDE4D Isoforms

FIG. 4

PDE4D isoform expression in EBV transformed cell lines (expression of PDE4D3 and -4D8 below detection limits).



PDE4D Isoforms

FIG. 5

Expression of PDE4D isoforms in EBV transformed cells from patients with or without the stroke associated haplotype

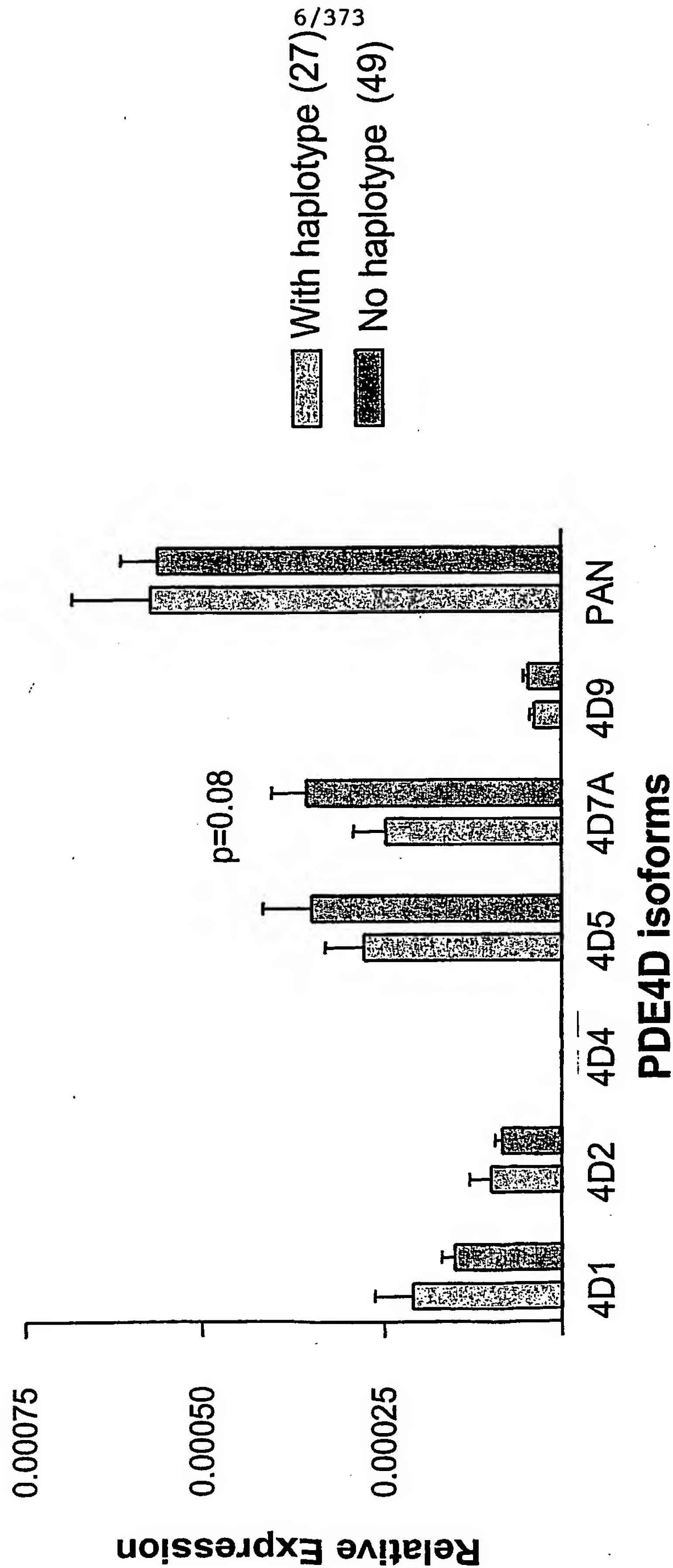


FIG. 6

Expression of PDE4D isoforms in EBV cells from controls with or without the stroke associated haplotype

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<210> 2
 <211> 809
 <212> PRT
 <213> Homo Sapien

<400> 2
 Met Glu Ala Glu Gly Ser Ser Ala Pro Ala Arg Ala Gly Ser Gly Glu
 1 5 10 15
 Gly Ser Asp Ser Ala Gly Gly Ala Thr Leu Lys Ala Pro Lys His Leu
 20 25 30
 Trp Arg His Glu Gln His His Gln Tyr Pro Leu Arg Gln Pro Gln Phe
 35 40 45
 Arg Leu Leu His Pro His His His Leu Pro Pro Pro Pro Pro Pro Ser
 50 55 60
 Pro Gln Pro Gln Pro Gln Cys Pro Leu Gln Pro Pro Pro Pro Pro Pro
 65 70 75 80
 Leu Pro Pro Pro Pro Pro Pro Pro Gly Ala Ala Arg Gly Arg Tyr Ala
 85 90 95
 Ser Ser Gly Ala Thr Gly Arg Val Arg His Arg Gly Tyr Ser Asp Thr
 100 105 110
 Glu Arg Tyr Leu Tyr Cys Arg Ala Met Asp Arg Thr Ser Tyr Ala Val
 115 120 125
 Glu Thr Gly His Arg Pro Gly Leu Lys Lys Ser Arg Met Ser Trp Pro
 130 135 140
 Ser Ser Phe Gln Gly Leu Arg Arg Phe Asp Val Asp Asn Gly Thr Ser
 145 150 155 160
 Ala Gly Arg Ser Pro Leu Asp Pro Met Thr Ser Pro Gly Ser Gly Leu
 165 170 175
 Ile Leu Gln Ala Asn Phe Val His Ser Gln Arg Arg Glu Ser Phe Leu
 180 185 190
 Tyr Arg Ser Asp Ser Asp Tyr Asp Leu Ser Pro Lys Ser Met Ser Arg
 195 200 205
 Asn Ser Ser Ile Ala Ser Asp Ile His Gly Asp Asp Leu Ile Val Thr
 210 215 220
 Pro Phe Ala Gln Val Leu Ala Ser Leu Arg Thr Val Arg Asn Asn Phe
 225 230 235 240
 Ala Ala Leu Thr Asn Leu Gln Asp Arg Ala Pro Ser Lys Arg Ser Pro
 245 250 255
 Met Cys Asn Gln Pro Ser Ile Asn Lys Ala Thr Ile Thr Glu Glu Ala
 260 265 270
 Tyr Gln Lys Leu Ala Ser Glu Thr Leu Glu Glu Leu Asp Trp Cys Leu
 275 280 285
 Asp Gln Leu Glu Thr Leu Gln Thr Arg His Ser Val Ser Glu Met Ala
 290 295 300
 Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His Leu Ser
 305 310 315 320
 Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser Asn Thr
 325 330 335
 Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr Gln Lys
 340 345 350
 Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly Val Lys
 355 360 365
 Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro Arg Phe
 370 375 380
 Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu Glu Asp
 385 390 395 400
 Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu Ser Gly
 405 410 415
 Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu Arg Asp
 420 425 430
 Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr Tyr Leu
 435 440 445
 Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His Asn Asn
 450 455 460
 Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu Ser Thr
 465 470 475 480
 Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala Ala Ile
 485 490 495

Fig. 7.1

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Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser Asn Gln
 500 505 510
 Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn Asp Ser
 515 520 525
 Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu Leu Gln
 530 535 540
 Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln Arg Gln
 545 550 555 560
 Ser Leu Arg Lys Met Val Ile Asp Ile Val Leu Ala Thr Asp Met Ser
 565 570 575
 Lys His Met Asn Leu Leu Ala Asp Leu Lys Thr Met Val Glu Thr Lys
 580 585 590
 Lys Val Thr Ser Ser Gly Val Leu Leu Leu Asp Asn Tyr Ser Asp Arg
 595 600 605
 Ile Gln Val Leu Gln Asn Met Val His Cys Ala Asp Leu Ser Asn Pro
 610 615 620
 Thr Lys Pro Leu Gln Leu Tyr Arg Gln Trp Thr Asp Arg Ile Met Glu
 625 630 635 640
 Glu Phe Phe Arg Gln Gly Asp Arg Glu Arg Glu Arg Gly Met Glu Ile
 645 650 655
 Ser Pro Met Cys Asp Lys His Asn Ala Ser Val Glu Lys Ser Gln Val
 660 665 670
 Gly Phe Ile Asp Tyr Ile Val His Pro Leu Trp Glu Thr Trp Ala Asp
 675 680 685
 Leu Val His Pro Asp Ala Gln Asp Ile Leu Asp Thr Leu Glu Asp Asn
 690 695 700
 Arg Glu Trp Tyr Gln Ser Thr Ile Pro Gln Ser Pro Ser Pro Ala Pro
 705 710 715 720
 Asp Asp Pro Glu Glu Gly Arg Gln Gly Gln Thr Glu Lys Phe Gln Phe
 725 730 735
 Glu Leu Thr Leu Glu Glu Asp Gly Glu Ser Asp Thr Glu Lys Asp Ser
 740 745 750
 Gly Ser Gln Val Glu Glu Asp Thr Ser Cys Ser Asp Ser Lys Thr Leu
 755 760 765
 Cys Thr Gln Asp Ser Glu Ser Thr Glu Ile Pro Leu Asp Glu Gln Val
 770 775 780
 Glu Glu Glu Ala Val Gly Glu Glu Glu Glu Ser Gln Pro Glu Ala Cys
 785 790 795 800
 Val Ile Asp Asp Arg Ser Pro Asp Thr
 805

<210> 3
 <211> 150
 <212> PRT
 <213> Homo Sapien

<400> 3
 Met Asp Arg Thr Ser Tyr Ala Val Glu Thr Gly His Arg Pro Gly Leu
 1 5 10 15
 Lys Lys Ser Arg Met Ser Trp Pro Ser Ser Phe Gln Gly Leu Arg Arg
 20 25 30
 Phe Asp Val Asp Asn Gly Thr Ser Ala Gly Arg Ser Pro Leu Asp Pro
 35 40 45
 Met Thr Ser Pro Gly Ser Gly Leu Ile Leu Gln Ala Asn Phe Val His
 50 55 60
 Ser Gln Arg Arg Glu Ser Phe Leu Tyr Arg Ser Asp Ser Asp Tyr Asp
 65 70 75 80
 Leu Ser Pro Lys Ser Met Ser Arg Asn Ser Ser Ile Ala Ser Asp Ile
 85 90 95
 His Gly Asp Asp Leu Ile Val Thr Pro Phe Ala Gln Val Leu Ala Ser
 100 105 110
 Leu Arg Thr Val Arg Asn Asn Phe Ala Ala Leu Thr Asn Leu Gln Asp
 115 120 125
 Arg Ala Pro Ser Lys Arg Ser Pro Met Cys Asn Gln Pro Ser Ile Asn
 130 135 140
 Lys Ala Thr Ile Thr Val

Fig. 7.2

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145

150

<210> 4

<211> 745

<212> PRT

<213> Homo Sapien

<400> 4

```

Met Ala Gln Gln Thr Ser Pro Asp Thr Leu Thr Val Pro Glu Val Asp
 1          5          10          15
Asn Pro His Cys Pro Asn Pro Trp Leu Asn Glu Asp Leu Val Lys Ser
          20          25          30
Leu Arg Glu Asn Leu Leu Gln His Glu Lys Ser Lys Thr Ala Arg Lys
          35          40          45
Ser Val Ser Pro Lys Leu Ser Pro Val Ile Ser Pro Arg Asn Ser Pro
          50          55          60
Arg Leu Leu Arg Arg Met Leu Leu Ser Ser Asn Ile Pro Lys Gln Arg
65          70          75          80
Arg Phe Thr Val Ala His Thr Cys Phe Asp Val Asp Asn Gly Thr Ser
          85          90          95
Ala Gly Arg Ser Pro Leu Asp Pro Met Thr Ser Pro Gly Ser Gly Leu
          100          105          110
Ile Leu Gln Ala Asn Phe Val His Ser Gln Arg Arg Glu Ser Phe Leu
          115          120          125
Tyr Arg Ser Asp Ser Asp Tyr Asp Leu Ser Pro Lys Ser Met Ser Arg
130          135          140
Asn Ser Ser Ile Ala Ser Asp Ile His Gly Asp Asp Leu Ile Val Thr
145          150          155          160
Pro Phe Ala Gln Val Leu Ala Ser Leu Arg Thr Val Arg Asn Asn Phe
          165          170          175
Ala Ala Leu Thr Asn Leu Gln Asp Arg Ala Pro Ser Lys Arg Ser Pro
          180          185          190
Met Cys Asn Gln Pro Ser Ile Asn Lys Ala Thr Ile Thr Glu Glu Ala
          195          200          205
Tyr Gln Lys Leu Ala Ser Glu Thr Leu Glu Glu Leu Asp Trp Cys Leu
210          215          220
Asp Gln Leu Glu Thr Leu Gln Thr Arg His Ser Val Ser Glu Met Ala
225          230          235          240
Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His Leu Ser
          245          250          255
Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser Asn Thr
          260          265          270
Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr Gln Lys
          275          280          285
Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly Val Lys
290          295          300
Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro Arg Phe
305          310          315          320
Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu Glu Asp
          325          330          335
Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu Ser Gly
          340          345          350
Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu Arg Asp
          355          360          365
Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr Tyr Leu
370          375          380
Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His Asn Asn
385          390          395          400
Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu Ser Thr
          405          410          415
Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala Ala Ile
          420          425          430
Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser Asn Gln
          435          440          445
Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn Asp Ser
450          455          460

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Fig. 7.3

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Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu Leu Gln
 465 470 475 480
 Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln Arg Gln
 485 490 495
 Ser Leu Arg Lys Met Val Ile Asp Ile Val Leu Ala Thr Asp Met Ser
 500 505 510
 Lys His Met Asn Leu Leu Ala Asp Leu Lys Thr Met Val Glu Thr Lys
 515 520 525
 Lys Val Thr Ser Ser Gly Val Leu Leu Leu Asp Asn Tyr Ser Asp Arg
 530 535 540
 Ile Gln Val Leu Gln Asn Met Val His Cys Ala Asp Leu Ser Asn Pro
 545 550 555 560
 Thr Lys Pro Leu Gln Leu Tyr Arg Gln Trp Thr Asp Arg Ile Met Glu
 565 570 575
 Glu Phe Phe Arg Gln Gly Asp Arg Glu Arg Glu Arg Gly Met Glu Ile
 580 585 590
 Ser Pro Met Cys Asp Lys His Asn Ala Ser Val Glu Lys Ser Gln Val
 595 600 605
 Gly Phe Ile Asp Tyr Ile Val His Pro Leu Trp Glu Thr Trp Ala Asp
 610 615 620
 Leu Val His Pro Asp Ala Gln Asp Ile Leu Asp Thr Leu Glu Asp Asn
 625 630 635 640
 Arg Glu Trp Tyr Gln Ser Thr Ile Pro Gln Ser Pro Ser Pro Ala Pro
 645 650 655
 Asp Asp Pro Glu Glu Gly Arg Gln Gly Gln Thr Glu Lys Phe Gln Phe
 660 665 670
 Glu Leu Thr Leu Glu Glu Asp Gly Glu Ser Asp Thr Glu Lys Asp Ser
 675 680 685
 Gly Ser Gln Val Glu Glu Asp Thr Ser Cys Ser Asp Ser Lys Thr Leu
 690 695 700
 Cys Thr Gln Asp Ser Glu Ser Thr Glu Ile Pro Leu Asp Glu Gln Val
 705 710 715 720
 Glu Glu Glu Ala Val Gly Glu Glu Glu Glu Ser Gln Pro Glu Ala Cys
 725 730 735
 Val Ile Asp Asp Arg Ser Pro Asp Thr
 740 745

<210> 5
 <211> 215
 <212> PRT
 <213> Homo Sapien

<400> 5

Met Ala Gln Gln Thr Ser Pro Asp Thr Leu Thr Val Pro Glu Val Asp
 1 5 10 15
 Asn Pro His Cys Pro Asn Pro Trp Leu Asn Glu Asp Leu Val Lys Ser
 20 25 30
 Leu Arg Glu Asn Leu Leu Gln His Glu Lys Ser Lys Thr Ala Arg Lys
 35 40 45
 Ser Val Ser Pro Lys Leu Ser Pro Val Ile Ser Pro Arg Asn Ser Pro
 50 55 60
 Arg Leu Leu Arg Arg Met Leu Leu Ser Ser Asn Ile Pro Lys Gln Arg
 65 70 75 80
 Arg Phe Thr Val Ala His Thr Cys Phe Asp Val Asp Asn Gly Thr Ser
 85 90 95
 Ala Gly Arg Ser Pro Leu Asp Pro Met Thr Ser Pro Gly Ser Gly Leu
 100 105 110
 Ile Leu Gln Ala Asn Phe Val His Ser Gln Arg Arg Glu Ser Phe Leu
 115 120 125
 Tyr Arg Ser Asp Ser Asp Tyr Asp Leu Ser Pro Lys Ser Met Ser Arg
 130 135 140
 Asn Ser Ser Ile Ala Ser Asp Ile His Gly Asp Asp Leu Ile Val Thr
 145 150 155 160
 Pro Phe Ala Gln Val Leu Ala Ser Leu Arg Thr Val Arg Asn Asn Phe

Fig. 7.4

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| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|--|
| | | | | 165 | | | | 170 | | | | | 175 | | | | |
| Ala | Ala | Leu | Thr | Asn | Leu | Gln | Asp | Arg | Ala | Pro | Ser | Lys | Arg | Ser | Pro | | |
| | | | 180 | | | | | 185 | | | | | 190 | | | | |
| Met | Cys | Asn | Gln | Pro | Ser | Ile | Asn | Lys | Ala | Thr | Ile | Thr | Gly | Leu | Tyr | | |
| | | 195 | | | | | 200 | | | | | 205 | | | | | |
| Asn | Gly | Ile | Ile | Ala | Phe | Leu | | | | | | | | | | | |
| | 210 | | | | | 215 | | | | | | | | | | | |

<210> 6
 <211> 673
 <212> PRT
 <213> Homo Sapien

<400> 6

| | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|--|
| Met | Met | His | Val | Asn | Asn | Phe | Pro | Phe | Arg | Arg | His | Ser | Trp | Ile | Cys | | |
| 1 | | | | 5 | | | | 10 | | | | | | 15 | | | |
| Phe | Asp | Val | Asp | Asn | Gly | Thr | Ser | Ala | Gly | Arg | Ser | Pro | Leu | Asp | Pro | | |
| | | 20 | | | | | | 25 | | | | | 30 | | | | |
| Met | Thr | Ser | Pro | Gly | Ser | Gly | Leu | Ile | Leu | Gln | Ala | Asn | Phe | Val | His | | |
| | | 35 | | | | | 40 | | | | | 45 | | | | | |
| Ser | Gln | Arg | Arg | Glu | Ser | Phe | Leu | Tyr | Arg | Ser | Asp | Ser | Asp | Tyr | Asp | | |
| | 50 | | | | | 55 | | | | | 60 | | | | | | |
| Leu | Ser | Pro | Lys | Ser | Met | Ser | Arg | Asn | Ser | Ser | Ile | Ala | Ser | Asp | Ile | | |
| 65 | | | | 70 | | | | 75 | | | | | | | 80 | | |
| His | Gly | Asp | Asp | Leu | Ile | Val | Thr | Pro | Phe | Ala | Gln | Val | Leu | Ala | Ser | | |
| | | | | 85 | | | | 90 | | | | | | 95 | | | |
| Leu | Arg | Thr | Val | Arg | Asn | Asn | Phe | Ala | Ala | Leu | Thr | Asn | Leu | Gln | Asp | | |
| | | | 100 | | | | | 105 | | | | | 110 | | | | |
| Arg | Ala | Pro | Ser | Lys | Arg | Ser | Pro | Met | Cys | Asn | Gln | Pro | Ser | Ile | Asn | | |
| | | 115 | | | | | 120 | | | | | 125 | | | | | |
| Lys | Ala | Thr | Ile | Thr | Glu | Glu | Ala | Tyr | Gln | Lys | Leu | Ala | Ser | Glu | Thr | | |
| | 130 | | | | | 135 | | | | | 140 | | | | | | |
| Leu | Glu | Glu | Leu | Asp | Trp | Cys | Leu | Asp | Gln | Leu | Glu | Thr | Leu | Gln | Thr | | |
| 145 | | | | 150 | | | | | | 155 | | | | | 160 | | |
| Arg | His | Ser | Val | Ser | Glu | Met | Ala | Ser | Asn | Lys | Phe | Lys | Arg | Met | Leu | | |
| | | | | 165 | | | | 170 | | | | | | 175 | | | |
| Asn | Arg | Glu | Leu | Thr | His | Leu | Ser | Glu | Met | Ser | Arg | Ser | Gly | Asn | Gln | | |
| | | | 180 | | | | | 185 | | | | | 190 | | | | |
| Val | Ser | Glu | Phe | Ile | Ser | Asn | Thr | Phe | Leu | Asp | Lys | Gln | His | Glu | Val | | |
| | | 195 | | | | | 200 | | | | | 205 | | | | | |
| Glu | Ile | Pro | Ser | Pro | Thr | Gln | Lys | Glu | Lys | Glu | Lys | Lys | Lys | Arg | Pro | | |
| | 210 | | | | | 215 | | | | | | 220 | | | | | |
| Met | Ser | Gln | Ile | Ser | Gly | Val | Lys | Lys | Leu | Met | His | Ser | Ser | Ser | Leu | | |
| 225 | | | | 230 | | | | 235 | | | | | | | 240 | | |
| Thr | Asn | Ser | Ser | Ile | Pro | Arg | Phe | Gly | Val | Lys | Thr | Glu | Gln | Glu | Asp | | |
| | | | | 245 | | | | 250 | | | | | | 255 | | | |
| Val | Leu | Ala | Lys | Glu | Leu | Glu | Asp | Val | Asn | Lys | Trp | Gly | Leu | His | Val | | |
| | | 260 | | | | | | 265 | | | | | 270 | | | | |
| Phe | Arg | Ile | Ala | Glu | Leu | Ser | Gly | Asn | Arg | Pro | Leu | Thr | Val | Ile | Met | | |
| | | 275 | | | | | 280 | | | | | 285 | | | | | |
| His | Thr | Ile | Phe | Gln | Glu | Arg | Asp | Leu | Leu | Lys | Thr | Phe | Lys | Ile | Pro | | |
| | 290 | | | | | 295 | | | | | 300 | | | | | | |
| Val | Asp | Thr | Leu | Ile | Thr | Tyr | Leu | Met | Thr | Leu | Glu | Asp | His | Tyr | His | | |
| 305 | | | | 310 | | | | | | 315 | | | | | 320 | | |
| Ala | Asp | Val | Ala | Tyr | His | Asn | Asn | Ile | His | Ala | Ala | Asp | Val | Val | Gln | | |
| | | | 325 | | | | | 330 | | | | | | 335 | | | |
| Ser | Thr | His | Val | Leu | Leu | Ser | Thr | Pro | Ala | Leu | Glu | Ala | Val | Phe | Thr | | |
| | | | 340 | | | | | 345 | | | | | 350 | | | | |
| Asp | Leu | Glu | Ile | Leu | Ala | Ala | Ile | Phe | Ala | Ser | Ala | Ile | His | Asp | Val | | |
| | | 355 | | | | | 360 | | | | | 365 | | | | | |
| Asp | His | Pro | Gly | Val | Ser | Asn | Gln | Phe | Leu | Ile | Asn | Thr | Asn | Ser | Glu | | |
| | 370 | | | | | 375 | | | | | 380 | | | | | | |
| Leu | Ala | Leu | Met | Tyr | Asn | Asp | Ser | Ser | Val | Leu | Glu | Asn | His | His | Leu | | |
| 385 | | | | 390 | | | | | | 395 | | | | | 400 | | |
| Ala | Val | Gly | Phe | Lys | Leu | Leu | Gln | Glu | Glu | Asn | Cys | Asp | Ile | Phe | Gln | | |
| | | | | 405 | | | | 410 | | | | | | 415 | | | |

Fig. 7.5

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Asn Leu Thr Lys Lys Gln Arg Gln Ser Leu Arg Lys Met Val Ile Asp
 420 425 430
 Ile Val Leu Ala Thr Asp Met Ser Lys His Met Asn Leu Leu Ala Asp
 435 440 445
 Leu Lys Thr Met Val Glu Thr Lys Lys Val Thr Ser Ser Gly Val Leu
 450 455 460
 Leu Leu Asp Asn Tyr Ser Asp Arg Ile Gln Val Leu Gln Asn Met Val
 465 470 475 480
 His Cys Ala Asp Leu Ser Asn Pro Thr Lys Pro Leu Gln Leu Tyr Arg
 485 490 495
 Gln Trp Thr Asp Arg Ile Met Glu Glu Phe Phe Arg Gln Gly Asp Arg
 500 505 510
 Glu Arg Glu Arg Gly Met Glu Ile Ser Pro Met Cys Asp Lys His Asn
 515 520 525
 Ala Ser Val Glu Lys Ser Gln Val Gly Phe Ile Asp Tyr Ile Val His
 530 535 540
 Pro Leu Trp Glu Thr Trp Ala Asp Leu Val His Pro Asp Ala Gln Asp
 545 550 555 560
 Ile Leu Asp Thr Leu Glu Asp Asn Arg Glu Trp Tyr Gln Ser Thr Ile
 565 570 575
 Pro Gln Ser Pro Ser Pro Ala Pro Asp Asp Pro Glu Glu Gly Arg Gln
 580 585 590
 Gly Gln Thr Glu Lys Phe Gln Phe Glu Leu Thr Leu Glu Glu Asp Gly
 595 600 605
 Glu Ser Asp Thr Glu Lys Asp Ser Gly Ser Gln Val Glu Glu Asp Thr
 610 615 620
 Ser Cys Ser Asp Ser Lys Thr Leu Cys Thr Gln Asp Ser Glu Ser Thr
 625 630 635 640
 Glu Ile Pro Leu Asp Glu Gln Val Glu Glu Ala Val Gly Glu Glu
 645 650 655
 Glu Glu Ser Gln Pro Glu Ala Cys Val Ile Asp Asp Arg Ser Pro Asp
 660 665 670
 Thr

<210> 7
 <211> 15
 <212> PRT
 <213> Homo Sapien

<400> 7
 Met Met His Val Asn Asn Phe Pro Phe Arg Arg His Ser Trp Ile
 1 5 10 15

<210> 8
 <211> 687
 <212> PRT
 <213> Homo Sapien

<400> 8
 Met Ala Phe Val Trp Asp Pro Leu Gly Ala Thr Val Pro Gly Pro Ser
 1 5 10 15
 Thr Arg Ala Lys Ser Arg Leu Arg Phe Ser Lys Ser Tyr Ser Phe Asp
 20 25 30
 Val Asp Asn Gly Thr Ser Ala Gly Arg Ser Pro Leu Asp Pro Met Thr
 35 40 45
 Ser Pro Gly Ser Gly Leu Ile Leu Gln Ala Asn Phe Val His Ser Gln
 50 55 60
 Arg Arg Glu Ser Phe Leu Tyr Arg Ser Asp Ser Asp Tyr Asp Leu Ser
 65 70 75 80
 Pro Lys Ser Met Ser Arg Asn Ser Ser Ile Ala Ser Asp Ile His Gly
 85 90 95
 Asp Asp Leu Ile Val Thr Pro Phe Ala Gln Val Leu Ala Ser Leu Arg
 100 105 110
 Thr Val Arg Asn Asn Phe Ala Ala Leu Thr Asn Leu Gln Asp Arg Ala

Fig. 7.6

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | 115 | | | | 120 | | | | | 125 | | | | |
| Pro | Ser | Lys | Arg | Ser | Pro | Met | Cys | Asn | Gln | Pro | Ser | Ile | Asn | Lys | Ala |
| | 130 | | | | | 135 | | | | | 140 | | | | |
| Thr | Ile | Thr | Glu | Glu | Ala | Tyr | Gln | Lys | Leu | Ala | Ser | Glu | Thr | Leu | Glu |
| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| Glu | Leu | Asp | Trp | Cys | Leu | Asp | Gln | Leu | Glu | Thr | Leu | Gln | Thr | Arg | His |
| | | | | 165 | | | | | 170 | | | | | 175 | |
| Ser | Val | Ser | Glu | Met | Ala | Ser | Asn | Lys | Phe | Lys | Arg | Met | Leu | Asn | Arg |
| | | | 180 | | | | | 185 | | | | | 190 | | |
| Glu | Leu | Thr | His | Leu | Ser | Glu | Met | Ser | Arg | Ser | Gly | Asn | Gln | Val | Ser |
| | | 195 | | | | | 200 | | | | | 205 | | | |
| Glu | Phe | Ile | Ser | Asn | Thr | Phe | Leu | Asp | Lys | Gln | His | Glu | Val | Glu | Ile |
| | 210 | | | | | 215 | | | | | 220 | | | | |
| Pro | Ser | Pro | Thr | Gln | Lys | Glu | Lys | Glu | Lys | Lys | Lys | Arg | Pro | Met | Ser |
| 225 | | | | | 230 | | | | | 235 | | | | | 240 |
| Gln | Ile | Ser | Gly | Val | Lys | Lys | Leu | Met | His | Ser | Ser | Ser | Leu | Thr | Asn |
| | | | | 245 | | | | | 250 | | | | | 255 | |
| Ser | Ser | Ile | Pro | Arg | Phe | Gly | Val | Lys | Thr | Glu | Gln | Glu | Asp | Val | Leu |
| | | | 260 | | | | | 265 | | | | | 270 | | |
| Ala | Lys | Glu | Leu | Glu | Asp | Val | Asn | Lys | Trp | Gly | Leu | His | Val | Phe | Arg |
| | | 275 | | | | | 280 | | | | | 285 | | | |
| Ile | Ala | Glu | Leu | Ser | Gly | Asn | Arg | Pro | Leu | Thr | Val | Ile | Met | His | Thr |
| | 290 | | | | | 295 | | | | | 300 | | | | |
| Ile | Phe | Gln | Glu | Arg | Asp | Leu | Leu | Lys | Thr | Phe | Lys | Ile | Pro | Val | Asp |
| 305 | | | | | 310 | | | | | 315 | | | | | 320 |
| Thr | Leu | Ile | Thr | Tyr | Leu | Met | Thr | Leu | Glu | Asp | His | Tyr | His | Ala | Asp |
| | | | | 325 | | | | | 330 | | | | | 335 | |
| Val | Ala | Tyr | His | Asn | Asn | Ile | His | Ala | Ala | Asp | Val | Val | Gln | Ser | Thr |
| | | | 340 | | | | | 345 | | | | | 350 | | |
| His | Val | Leu | Leu | Ser | Thr | Pro | Ala | Leu | Glu | Ala | Val | Phe | Thr | Asp | Leu |
| | | 355 | | | | | 360 | | | | | 365 | | | |
| Glu | Ile | Leu | Ala | Ala | Ile | Phe | Ala | Ser | Ala | Ile | His | Asp | Val | Asp | His |
| | 370 | | | | | 375 | | | | | 380 | | | | |
| Pro | Gly | Val | Ser | Asn | Gln | Phe | Leu | Ile | Asn | Thr | Asn | Ser | Glu | Leu | Ala |
| 385 | | | | | 390 | | | | | 395 | | | | | 400 |
| Leu | Met | Tyr | Asn | Asp | Ser | Ser | Val | Leu | Glu | Asn | His | His | Leu | Ala | Val |
| | | | | 405 | | | | | 410 | | | | | 415 | |
| Gly | Phe | Lys | Leu | Leu | Gln | Glu | Glu | Asn | Cys | Asp | Ile | Phe | Gln | Asn | Leu |
| | | | 420 | | | | | 425 | | | | | 430 | | |
| Thr | Lys | Lys | Gln | Arg | Gln | Ser | Leu | Arg | Lys | Met | Val | Ile | Asp | Ile | Val |
| | | 435 | | | | | 440 | | | | | 445 | | | |
| Leu | Ala | Thr | Asp | Met | Ser | Lys | His | Met | Asn | Leu | Leu | Ala | Asp | Leu | Lys |
| | 450 | | | | | 455 | | | | | 460 | | | | |
| Thr | Met | Val | Glu | Thr | Lys | Lys | Val | Thr | Ser | Ser | Gly | Val | Leu | Leu | Leu |
| 465 | | </ | | | | | | | | | | | | | |

Fig. 7.7

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Pro Leu Asp Glu Gln Val Glu Glu Glu Ala Val Gly Glu Glu Glu Glu
 660 665 670
 Ser Gln Pro Glu Ala Cys Val Ile Asp Asp Arg Ser Pro Asp Thr
 675 680 685

<210> 9
 <211> 585
 <212> PRT
 <213> Homo Sapien

<400> 9
 Met Lys Glu Gln Pro Ser Cys Ala Gly Thr Gly His Pro Ser Met Ala
 1 5 10 15
 Gly Tyr Gly Arg Met Ala Pro Phe Glu Leu Ala Ser Gly Pro Val Lys
 20 25 30
 Arg Leu Arg Thr Glu Ser Pro Phe Pro Cys Leu Phe Ala Glu Glu Ala
 35 40 45
 Tyr Gln Lys Leu Ala Ser Glu Thr Leu Glu Glu Leu Asp Trp Cys Leu
 50 55 60
 Asp Gln Leu Glu Thr Leu Gln Thr Arg His Ser Val Ser Glu Met Ala
 65 70 75 80
 Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His Leu Ser
 85 90 95
 Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser Asn Thr
 100 105 110
 Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr Gln Lys
 115 120 125
 Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly Val Lys
 130 135 140
 Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro Arg Phe
 145 150 155 160
 Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu Glu Asp
 165 170 175
 Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu Ser Gly
 180 185 190
 Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu Arg Asp
 195 200 205
 Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr Tyr Leu
 210 215 220
 Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His Asn Asn
 225 230 235 240
 Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu Ser Thr
 245 250 255
 Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala Ala Ile
 260 265 270
 Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser Asn Gln
 275 280 285
 Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn Asp Ser
 290 295 300
 Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu Leu Gln
 305 310 315 320
 Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln Arg Gln
 325 330 335
 Ser Leu Arg Lys Met Val Ile Asp Ile Val Leu Ala Thr Asp Met Ser
 340 345 350
 Lys His Met Asn Leu Leu Ala Asp Leu Lys Thr Met Val Glu Thr Lys
 355 360 365
 Lys Val Thr Ser Ser Gly Val Leu Leu Leu Asp Asn Tyr Ser Asp Arg
 370 375 380
 Ile Gln Val Leu Gln Asn Met Val His Cys Ala Asp Leu Ser Asn Pro
 385 390 395 400
 Thr Lys Pro Leu Gln Leu Tyr Arg Gln Trp Thr Asp Arg Ile Met Glu
 405 410 415
 Glu Phe Phe Arg Gln Gly Asp Arg Glu Arg Glu Arg Gly Met Glu Ile
 420 425 430
 Ser Pro Met Cys Asp Lys His Asn Ala Ser Val Glu Lys Ser Gln Val

Fig. 7.8

15/373

435 440 445
 Gly Phe Ile Asp Tyr Ile Val His Pro Leu Trp Glu Thr Trp Ala Asp
 450 455 460
 Leu Val His Pro Asp Ala Gln Asp Ile Leu Asp Thr Leu Glu Asp Asn
 465 470 475 480
 Arg Glu Trp Tyr Gln Ser Thr Ile Pro Gln Ser Pro Ser Pro Ala Pro
 485 490 495
 Asp Asp Pro Glu Glu Gly Arg Gln Gly Gln Thr Glu Lys Phe Gln Phe
 500 505 510
 Glu Leu Thr Leu Glu Glu Asp Gly Glu Ser Asp Thr Glu Lys Asp Ser
 515 520 525
 Gly Ser Gln Val Glu Glu Asp Thr Ser Cys Ser Asp Ser Lys Thr Leu
 530 535 540
 Cys Thr Gln Asp Ser Glu Ser Thr Glu Ile Pro Leu Asp Glu Gln Val
 545 550 555 560
 Glu Glu Glu Ala Val Gly Glu Glu Glu Glu Ser Gln Pro Glu Ala Cys
 565 570 575
 Val Ile Asp Asp Arg Ser Pro Asp Thr
 580 585

<210> 10
 <211> 507
 <212> PRT
 <213> Homo Sapien

<400> 10
 Met Ala Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His
 1 5 10 15
 Leu Ser Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser
 20 25 30
 Asn Thr Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr
 35 40 45
 Gln Lys Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly
 50 55 60
 Val Lys Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro
 65 70 75 80
 Arg Phe Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu
 85 90 95
 Glu Asp Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu
 100 105 110
 Ser Gly Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu
 115 120 125
 Arg Asp Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr
 130 135 140
 Tyr Leu Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His
 145 150 155 160
 Asn Asn Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu
 165 170 175
 Ser Thr Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala
 180 185 190
 Ala Ile Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser
 195 200 205
 Asn Gln Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn
 210 215 220
 Asp Ser Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu
 225 230 235 240
 Leu Gln Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln
 245 250 255
 Arg Gln Ser Leu Arg Lys Met Val Ile Asp Ile Val Leu Ala Thr Asp
 260 265 270
 Met Ser Lys His Met Asn Leu Leu Ala Asp Leu Lys Thr Met Val Glu
 275 280 285
 Thr Lys Lys Val Thr Ser Ser Gly Val Leu Leu Leu Asp Asn Tyr Ser
 290 295 300
 Asp Arg Ile Gln Val Leu Gln Asn Met Val His Cys Ala Asp Leu Ser
 305 310 315 320

Fig. 7.9

16/373

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asn | Pro | Thr | Lys | Pro | Leu | Gln | Leu | Tyr | Arg | Gln | Trp | Thr | Asp | Arg | Ile |
| | | | | 325 | | | | | 330 | | | | | 335 | |
| Met | Glu | Glu | Phe | Phe | Arg | Gln | Gly | Asp | Arg | Glu | Arg | Glu | Arg | Gly | Met |
| | | | 340 | | | | | 345 | | | | | 350 | | |
| Glu | Ile | Ser | Pro | Met | Cys | Asp | Lys | His | Asn | Ala | Ser | Val | Glu | Lys | Ser |
| | | 355 | | | | | 360 | | | | | 365 | | | |
| Gln | Val | Gly | Phe | Ile | Asp | Tyr | Ile | Val | His | Pro | Leu | Trp | Glu | Thr | Trp |
| | 370 | | | | | 375 | | | | | 380 | | | | |
| Ala | Asp | Leu | Val | His | Pro | Asp | Ala | Gln | Asp | Ile | Leu | Asp | Thr | Leu | Glu |
| 385 | | | | | 390 | | | | | 395 | | | | | 400 |
| Asp | Asn | Arg | Glu | Trp | Tyr | Gln | Ser | Thr | Ile | Pro | Gln | Ser | Pro | Ser | Pro |
| | | | 405 | | | | | | 410 | | | | | 415 | |
| Ala | Pro | Asp | Asp | Pro | Glu | Glu | Gly | Arg | Gln | Gly | Gln | Thr | Glu | Lys | Phe |
| | | | 420 | | | | | 425 | | | | | 430 | | |
| Gln | Phe | Glu | Leu | Thr | Leu | Glu | Glu | Asp | Gly | Glu | Ser | Asp | Thr | Glu | Lys |
| | | 435 | | | | | 440 | | | | | 445 | | | |
| Asp | Ser | Gly | Ser | Gln | Val | Glu | Glu | Asp | Thr | Ser | Cys | Ser | Asp | Ser | Lys |
| | 450 | | | | | 455 | | | | | 460 | | | | |
| Thr | Leu | Cys | Thr | Gln | Asp | Ser | Glu | Ser | Thr | Glu | Ile | Pro | Leu | Asp | Glu |
| 465 | | | | | 470 | | | | | 475 | | | | | 480 |
| Gln | Val | Glu | Glu | Glu | Ala | Val | Gly | Glu | Glu | Glu | Ser | Gln | Pro | Glu | |
| | | | | 485 | | | | 490 | | | | | 495 | | |
| Ala | Cys | Val | Ile | Asp | Asp | Arg | Ser | Pro | Asp | Thr | | | | | |
| | | | 500 | | | | | 505 | | | | | | | |

Fig. 7.10

| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
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TGGTTGGATTGTTTATGCTCTTTTTTATTATTTATTCTTATTTTACCATGAAAATATCACTAAGTTCTTTGGTTTGTG
ACCTGATTGTACCTACTTTGACAAATCACTGCCTTTCTGGACCCAGTTTCTCATTAAGTGGCAGTGATAACCTGTCAT
ACTTACAGATATAAAAACATGAAAGTTAAAGTATTGGGTAATACTTTCTCTCTATCTTTTTTTTTTATTTTGAAAAGATA
AAAAATTGGCATAATGTATTAGTTAAGATGGAATAATCATATGTTGATATCCAGCCATTTCTTCTCTCAATGATAGGA
AGATTTTTATGTGAAACTACTTGTGAGAGATCTTAACAATTTGTAGTTAGAGAAAGCACTATTATATCATTTTGAAATG
CAAGAAACAAGTTACCTTTGGGGCAACAGAGGCCCTTGTCATTTTCTCAAAAGAAGGAAGCATCAGCATTTTGATGATG
ATGTTGAGATTGTAGAAATGATGAAGGTGAAAAAGTTATCTAGCTTATGTTTAGCAAAATGAAATGAACCCAAATAAT
AAAACAGTTACAACATTGAATCTCTTTGGGAGAAAAAAAAGATAGAATGCTAATGTCCTTCAGAACTTCTTAAACCA
GAACCTTAAAAAAAAGAGAAGCTTTTAAAAAATCATAATAGTTTATGATCTTGAAGGGTTTAAAAGTATTTGATGAAGA
TGTCTTTTGAATTTATTTGTAGGTCTTCTTGTGTATTTTAAAGCTAAGTTATCTTGTAATCATTTTTTTTCTATACCTTT
GTCAGTAACCTCTTAGTGATGAAATAAAAAAGATTAGGTAATCATCCAGCAATGGGGAAGAAGTTAAGGAACAAAGAGC
TCAGATTAACTAGTTTTTAGAATCTAAGCATTCTGCATGAATTTGAATCATGGAACAAATGTAGCACTCCAACA
TTTGATGCAAACTAAAAGTGGAATACTGCTTTGATATTTGAATGAATTGAAAAATAATTAACATCCTTGGAACCTGTAT
GTAAAGAAGGACTTCACAAGTATTATAGATACCCCCAACCTCAGCCCTTTTCCCATGTATCTCTTTGATCACATCCCTA
CCTCATAGATCACCCATGTGCTGAAGACTTTCAGTTCTGTATCTTCATTCTAGATCTCCTGAACCTCAAGATCAGAATAT
CTTTCTGACTTCTGACTGTGTATTTCTGGATGTTATACAAGAACCTCAGCTCAAACCTCAGTATTCCCTAAACCATTGTT
TTTGAAACTTTATGTTGGATGTGAAATCTGTATTGTAGAATAACATTAAAAAAAGAAAGATAGTATGCAAAATATCAG
AGTGCATTGTATGTAGCAAGAGTAGGTATTTTCGTAACTTTTTGTTTTAATTAAACACATATATATTTATTATTGCAG
TGTAATATGATTTCTTAATTTGGATAAGTGTTAGTGAGGATGTGGATAAATTGGAACCTTTGTACATTACTGGTGGGA
CTATAAATGGCACTGCCGTTTGGTAAAAACAGTTTGGCAGTTCTCAAAAAGTTAAACATACAGTTAACATGTGATATA
GAAATTTCACTTTTAGATGTACACCCAAAAGAATTGAGAACATATGTTACACAGCAACTTGACACAAATGTTTCATAG
CAGCATTACTCAGAAGAGCCAAAAAGTGGAACAACCTGAAATGTCCATCAAGTGATGAAGCAGTAAATGTAGTATATC
CGTACAATGAAATATTAGCCATAAAAAGGAATGCAATGTTGTTGCATGCTACAACAACCTGGATGAATCTTGGAACA
TTATTCTAAGTAAAAGATTCCATTTTTATGAAATGTCCAGAATAGGCAAATCTATAGAGACAAAGATAAGTGGTTTCCA
GGGGTTGTGGGGAGGAGAGAATGGGAAGGTGACAAAATGTTCTGGATTAGATAATAGGGATGGGTATAACTTAGTGACT
ATACAAAAAAATCACTAGAATCATATACTTTAAAAAAGATATTTCCATAAAAAAAGAACAAAGCAAGAAAAAATAACT
AAATTTGACTTTAGGAGTTAAAAAGAATATAGTATCTCAAATGAAAATTTTGCTGGATAGGATTAGGGGTAGATTAGAC
ACTCCAGAAGTTAAAGATCAGTGAGCTTGAATACACACAATAGAAGCTAGTCTAAACAAAGCACAGAGAGAAAAAGAA
CAAAACAAACCTCCCAACAACAAACAAACAAAGCCAACCAAAAATACAGCCTCAATGACCTATGGGAAATATTTAGC
AGTCTAATATACATGTAATTGGAATCCCTGAAGGAGGAGGGGGGTAGAATGTATCTTTTTTTTGTCCCCTATGACTGCTG
TTAAGATTTTATTATTGATTTTATAGGAATTGCATTATATCTTGGTGTGGTTGTTTAAACAGAGGTATAGCTTATCAACC
AATGGTGGAGCTAAAATAGAATACTTGAAAGTACTTATGGATGCACAGAATCTAAGATGGCCCCCAATTTTCTCTGCTAC

Fig. 9.1

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CTTGTACCCTTGAGTATATGTGGGACCTGTTACTTGCTTCTAACCAATAAAATCTCACACCAGTTAGAATGGTGATTAT
TAAAAAGTCAGGAAACAACGGATGCTGGAGAGGATGTGGAAAAATAGGAACGCTTTTACACTGTTGGTGGAAGTGTA
TTAGCTCAGCCATTGTGGAAGACAGTGGCAATTCCTCAAGGATCTAGAAGTACCAATTTGACCCAGCCATCCCG
TTACTGGGTATATAACCAGAGGATTATAAATCATTCTACTATAAAGACACATGCACACGTATGTTTATTGCGGCACTGT
TCACAATAGCAAAGACTCGGAACCAACCCAAATGTCCATCAGTGATAGACTGGATGAAGACAATGTAGCACATATACAC
CAGGGAATACTATGCAGCTATAAAAAATGACGAGTTCATGTCCTTTGCAGGGACATGGATGAAGCTGGAAACCATCATT
CTCAGCAAACCTATCACAAGAACAGAAAACCAAACACCACATCTTCTCACTCATAAGTGGCAGTTGAACAATGAGAACAC
ATGGACACAAGGCAGGGAACATCACACACCCGGGGCCTGTGTGGGGGGTCCGGGGGAGGGGATAGCATTAGGAGA
AATACCTAATGTAGTTGACATTACTTTGGTTTGGACATTACTTTGGTTTGTGGGTGCCACAAACCACCATGGCACATGTA
TACCTGTGTTACAAACCTGCACGTTCTGTACATGTACCCAGAACTTAAAGTATAATAATAATAAAATAACATGTATGT
CAAGGGTGACATGTAATTAAGCAAAGCTCAGTAAATTTAAATGATTGAAATTGTACTAAGTTTTCTGACCACGCTAGA
ATTAAGCTAGAACTCCAGGTCCAGATGGCTTACTGATCAATTTTACCCAAACACTTTGGAATGAATAATGACATTTGTAT
GAAAGTCCTTTCAGAAAATAGAAGATGAGGGAATATTTCTGAACCATTTTATGAGGCCAGTATTGACATGGGTAATAA
AACCAACAAATACATTACACAAAAAATTTGTAGCACATGATATCCCTGATAAAACCAAATGCAAAAATACATTAAATTT
GCAAATTGAATGCAGCAGTAGATAAAAAAGGACAATAATACATCATGGCCAAGTAGGGTTTATCCCAGCAAGGTAAGACT
GGTTTAACATCTAAATCAATCAGTATAATTCATCATATCGATAGGATGAAGGAAAAAACTCATGTGACCATCTCAAC
GATTGCAGAAAATGTATGTGACAATATTCACACCCATTAATGATAAAAAATGTTAAATACATTACAATAGAAGAAAAC
TCCTCAGCCTTATCAAGGGTACCTGTGAGAAAATTTATGGATAACATTTTCTTAATGGTGGTAGACTGAATGCTTTCCC
CTATGGTCAGAAAAAGACAAAACCTCATCACTGCTATAACAATTTTCATGAGAGGTCAGCAGTGCTTTCATGCCTTAAAG
GCATGAAAATGAAATAAGTGATTTAAGATTGGAAAGAAGAACTAAAACCTACGTTTGCTGATATCAAAAATCCCAAGAAA
TCTGCCCCCAAAAAGCACTTATGAATTAATAATTAACTTAACAAGGAAGCAGGATATAAGACCACTGTATAAAAATCA
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CCTGTTTTTAAGACTTACTATAAAATCTTACTTTCAGGTGTGGTATTGGTATCTTACTGTAAAGTCTTCTGTAAAGTA
TATTGATATTTAGTGTGGTGTGGCATAAGGATAGATATTTAGGCCATGGAATAGAATAGAGGGTCCAATAGTAGATT
CATGTATCTGTAGTCAAGTGATTTTCAGCAAAGAAGCCAAGGGAAGGGATCATCTTTTCAGGGTAGTGTGGAACAAC
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CCATGCAATGAAATATTACCCAGCAATAAAAATAAACTATGGGTACATAACAATATAAATGAATCTCAAAAATATGC
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GGTAATAAAGAAGTCAAAGCACTATTTGTGAAAATCAGTATATCATATGACGGTAAGCATAGTTGCTATTCACCAAAA
ACGTTTCAGAAAACATTTGAATTCATTGTCTGAAAGAGCTTAGGCTCAAGACTTGAATTACTAAGAAAAGAAAGTAGTAT
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CCCCACTCCCCCTACCCAGGACAGGCCCCGGTGTGTTATATTTCCCTTTCTGTGTTCAAGTGTCTCATTTGTTCAATG
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CTACAAAGGACATGAACTCATCTTTTTTATGGCTGCATAGTATTCATGGTGTATGTGTGCCACATTTTATTTTTTATT
TTTATTATTTTTTATTTTTTAAATTTTATTATATACTTTAAGTTAGTGATGTGCACAACATGCAGGTTTGTACAT
ATGTATATATGTGCCATGTTGGTGTGCTGCACCCATTAAGTCGTCATTTAACATTAGATATATCTCCTAATGCTATCCC
TCCCCCTACCCCGACCCCAACAGTCCCCGGTGTGTGATGTTCCCTTCTGTGTCAATGTGTTCTCATTTGTTCAATT
CCCACCTATGAGTGGCAACATGTGGTGTGTTGGTTTTTTGTCCCTGAGATAGTTTGTGAGAATGATGGTTTCCAGTTTC
ATCCATGTCCCTACAAAGCACATGAACTCATTTTTTTCATGGCTGCATAGTATTCCTGGTGTATAGTGCCACATTTT
CTTAATCCAGTCTATCACTGATGGACATTTGGGTGTTTCCAAGTCTTTGCTATTGTGAATAGTGCCTCAATAAACATA
CGTGTGCATGTGTCTTTATAGCAGCATGATTTATAATCCTTTGTGTATATACCCAGTAATGGGATGGCTGGGTCAAATG
GTATTTCTAGTTCTAGATCCTTGAGGAATCGCCACACTGTCTTCCACAATGGTTGAACCAGTTTACAGTCCCACCAACA

Fig. 9.2

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GTGTAAAAGCATTTCCTATTTCTCCACATCCTCTCCAGCACCTGTTGTTTCCTGACTTTTTTAATGATCGCCATTCTAACT
GGTGTGAGATGGTATCTCATTGTGGTTTTGATTTGCATTTCTCTGATGGCCAGTGATGGTGAGCATTTTTTTCATGTGTC
TTTTGGCTGCATAAATGTCTTCTTTTGAGAAGTGTCTGTTTCATATCCTTCGCCTACTTGTTGATGGGGTTGTTTGT
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TCCCATCCTGTAGGTTGCTTGTTCACCTCTGATGGTAGTTTCTTTTGCTGTGCAGAAGCTCTTTAGTTTAAATTAGATCCT
GTTTGTCAATTTTGGCTTTTGTTCCTTGTCTTTTGGTGTTTTAGACATGAAGTCCTTGCCCATGCCTATGTCCTGAATG
GTATTGCCTAGGTTTTCTTCTAGGGTTTTATGGTTTTAGGTCTAGCATTTAAGTCTTTAATCCATCTTGAATTAATTT
TTGTATAAGGTGTAAGGAAGGGATCTAGTTTCAGCTTTCTACATATGGCTAGCCAGTTTTCCAGCACCATTATTTAA
TAGGGAATCATTTCCTCATTTCTTGTTTTTGTTCAGGTTTGTCAAAGATCAGGTAGTTGTAGATATGTGGCATTATTTCT
GAGGGCTCTGTTCTGTTCCATTGGTCTATATCTCTGTTTTGGTACCAGTACCATGCTGTTTTGGTGACTGTAGCCTTGT
AGTATAGTTTGAAGTCAGGTACCGTGATGCCTCCAGCTTTGTTCTTTTGGCTTAGGATTGATTGGTAATGCGGGCTCT
TGTTTGGTTCATATGAACCTTAAAGTAGTTTTTCCAATTCTGTAAAGAAAGTCATTGGTAGCTTGATGGGGATGGCA
TTGAATCTATAAATTACCTTGGGCAGTATGGCCATTTTCAACAATATTGATTCTTCTACCCATGAGCATGGAATGTTCC
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GCTTAAGGAGATTTTGGGCTGAGACGATGGGGTTTTCTAGATATACAATCATGTCTGCAAACAGGGACAATTTGAC
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GATTATGGTGGATCAGCTTTTTGATGTGCTGCTGGATTCCGGTTTGCCAGTACTTTATTGAGGATTTGTTTCATTGATGTA
CATCAGGGATATTAGTGTAATAATCTCTTTTTTTTGTGTGTCTCTGCCAGGCTTTGGTATCAGGATGATGCTGGCCTCA
TCAAATGAGTTAGGGAGGATTCCTCTTTTTTCTATTGATTGGAATAGTTTCAGAAGGAATGGTACCAGCTCCTCCTTGT
ACCTGTGGTAGAATTTGGCTGTGAATCCGTCTGGTCTGGACTTTTATTGGTTGGTAAGCTATTAATTATTGCCTCAAT
TTCAGAGCCTGTTATTGGTCTATTCAAGATTCAACTTCTTCTGGTTTAGTTTTGGGAGAGTGTATGTATCGAGGAAT
TTATCCATTTCTTCCAGATTTTCTAGTTTCATTTTCATAGAGGTGTTTATAGTATTCTCTGATGGTAGTTTGTATTTCTG
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TGCTAGCAGTCTATCAATTTTGTGATCTTTTCAAGAAACCAGCTCCTGGATTCAATTGATTTTTTGAAGGGTTTTTGT
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TATTGTGTGGGAGTCTACGTCTCTTAGTAGGTCTCTAAGGACTTGCTTTATGAATCTGGCTGCTCCTGTATTGGGTGCA
TATATATTTAGGATAGTTAGCTCTTCTTGTGTAATTGATCCCTTTATCATTTATGTAATGGCCTTCTTTGTCTCTTTTGA
CCTTTGTTGGTTTTAAAGTCTGTTTTATCAGAGACTAGGATTGCAACCCCTGCCTTATTTTGTTTTTCCATTTGCTTCGCA
GATCTTCTCCATCCCTTTAATTTGAGCCTATGTGTGTCTCTGCATGTGAGATGGGTTTCTGAATACAGCACACTGAT
GAGTCTTGACTCTTTATCCAATTTGCCAGTTTGGGTCTTTAATTGGAGCATTTAGCCCATTTACATTTAAGGTTAATA
TTGTTACGTGTGAATTTGATCCTGTCTATTATGATGTTAGCTGGTTAATTTGCCCTGTTAGTTGATGCAGTTTCTTCTTAG
CCTCGATGGTCTTTACAATTTGGCATGATTTTGCAGTGGCTGGTACCGTATGTTTCTTTCCATGTTTAGTGCTTCTCTC
AGGAGCTCTTTTAGGGCAGGCCTGGTGGTGACAAAATCTCTCAGCATTTGCTTGTCTGTAAAGGATTTTATTTCTCCTT
CACTTATGAAGCTTAGTTTTGGCTGGATATGAAATCTGGGTTGAAAATCTTTCTTTCAGGAATGTTGAATATTGGTCC
CCACTCTCTTCCGGCTTGTAGGGTTTTCTGCCGAGAGATCAGCTGTTAGTCTAATGGGCTTCCCTTTGTGGGTAACCTGA
CCTTTCTCTCTGGCTGCCCTTAACATTTTTTCTTTCATTTCAACTTTGGTGAATCTGACAATTATGTGTCTTGGAGTTG
CTCTTCTTGAGGAGTATCTTTGTGGCGTTCTCTGTATTTCTTGAATTTGAATGTTGGCCTGCCTTGCTAGATTGGAGAA
GTTCTCCTGGATAATATCCTGAAGAGTTTTTTTCCAACCTGGTTCCATTTCTCCCCGTCACTTTCAGGTACACCAATCAGA
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TCTCGCTTCATTTCAATTCATTTGATCTTCCATCACTGATAACCTTCCCTTCCATTTGATCGAATCAGCAACTGAGGCTTG
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TAGTTTGATCATCTGAAGCCTTCTGCTCTCAACTCGTCAAAGTCATTCTCCATCCAGCTTTGTTCTGTTGCTGCTGAGG
AGCTGCGTTCTTTGGAGGAGGAGAGGTGCTCTGATTTTATAGAGTTTCCAGTTTTTCTGCTCTGTTTTTCCCCCATCTT
TGTGGTTTTATTTACCTTTGGTCTTTGATGATGGTGACGTACAGATGGGGTTTTGGTGTGGATGTCCTTTCTGTTTGT
AGTTTTCTTCTAACAGTTAGGACCCTCAGCTGCAGGTCTGTTGGTGTGTTGCTGGAGGTCCACTCCAGACCCTGTTTGC
CTGGGTATCAGCAGCAGAGGCTGCAGAACAGCAGATATTGGTGAACAGCAAATGTTGCTGCCTGATCGTTCTTTTGGA
GTTTTTTCTCAGAGGAGTACCCGGCCATGTGAGGTGTCAATTCAGCCCCCTACTGCGGGGTGCCTCCAGTTAAGCTACTC

Fig. 9.3

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GGGAGTCAGGGACCCACTTGAGGAGGCAGTCTGTCCATTCTCAGATCTCAAGCTGCATGCTGGGAGAACCCTACTCTC
TTCAAGGCTGTCAGACAGTGACATTTAAGTCTGCAGAGGTATATTGCTCCCTTTTGTGGCTATGCCCTGCCCCCAGAG
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AAAGGGAATTCCTGACCCCTTGCTGCTTCCCAGGTGAGGCGATGCGTCACCATTTCTTTGGCTCACGCTTGGTGTGCTGC
ACCCACTGTCCTGCACCCACTGTCGACACTCCCCAGTGAGATGAACCCGTTATGTCAGTTGGAAATGCAGAAATCACC
CGTCTTCTGCGTTGCTCACGCTGGGCACTGTAGACTGGAGCTGTTCCCTATTTGGCCATCTTGGTTCCATCCCCCTACT
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CTTGGCTGCCCCAAGTGCTGGGATTACAGGCATCAGCCACCATGCCTGGCTGCTAATAATAACTTTAAAAAACCTAAC
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CAAATATAAACAACTTTACAAAACCTGAATGGATTAGCAGGGTGTGGGGCACATGCTTGTAGTTCCATCTCCTTGGA
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ACAAAACAGAGACAATTGGAGGAAGACAGAACTAGAAAGAAAGTGTAATAAATAAATAAACACCCCAAACTCACAGAA
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CAAGGCTCAGATGGGGAAACGCATGGTGCTGCTTTTAGAACCATGCTCAGAAAGATTTGGTGGAGTTTAAACAAACAAG
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GGAGTGCAGCGGCACAATCTTGGCTCATTTGCAACCTCCACTTCCTGGGCTCAAGCAATCTCCTGCCTCAGCCTCCCAA
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GCCAGGCTGGACTTGAGCTCCAGACCTCAGGTGATCCACTCACCTCAGCCTCCCAAAGTGCTAGGATTACAGGCATAAG
CCACTGCACCCAGCCAGATGTAGCATAATTCTTAAGGGCCCTAGGATTTTTTGAATGGTAAAGGAGCACTGGTTTCAAC
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CCCTTCTAGTTACAAATGTCCTAGATGGCATCTTCTCCCATTAAGGCTGTTTTTTGTCTACATTGGAAATCTGTTGTTT
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TTCTGTAGTTTCCCCACCTCCCTCAGCCTTTATAGAATTGAAGAGTTAGGACTTTTCTCTAGGTTAGGGTGGGGCTTAA
AGAAATGTTGTGATTGGTTGGATCTTCTATCTAGGCCACTCAAACCTTCTCCCTATCAGCAACACAGCTGTTTCACTGC
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GGCACCAGAGGCCTAGCTTGTGACTTCTCTCAGCTTTTGACCTGCCACCCTTACTAAGGTCAATAGTTTCTTTTGATTT
AAGGTGACAGATGTGTGACTCTTCTTCACTTGAACACTTAGAGGCCATTGTAGGGTTATTAATTGGCCCAATTTCAAT
ATTGTTGTGTCTGAGGGAATAGAGAGGCCCAAAGGGAGGGAGAGAGACAGGGGAACCACTGTTGGTGGAGCAGTCAGA
ACACACACATTGGTTGATTAAAGTTCACAGTCTTATGGGCATATTGTGTGGTTCCCCCAAACACTTACAGTAGTAACAGC
AAAGATTACTTATTGATCATAGGTCATAATAATAGATAAAATAATAATTAATAAATTTGAAATATTCTGAATTACCAAAA
TGTGATACAGAGACATGGTGTGAGCCCATGTTGTTGGAAAAATGGTGGTGATAGCCTTGATTAAACACAGGGTTGCCACA

Fig. 9.4

AAACTTCAATTTGTAAAAACATAATACCTGCAAAGCAACTAAAGTGAAGTGCAGTAAAACGAGGTTATACCTGTATAT
TAATAGGTGACTCCAATAAAGACTTCGGTAATCTATAACAAGGAGCCAACTATCAAATGGCAACTGCAAAGATAGTTCT
CTCACTGAAGCTAACAAAAACATCTACAACTTTTCAGCTGAAAAATCAAAAAAGTTTGAGTTGTATAGGACATTCTAAC
ACCAGGGAATGAGACATATCTTTTGTATGTAATAATAATGCAAGCCTGAAAGTCTTCCAGTGACTCACAGAGTAATAA
CTGTGACAGAGGCTTTCTGAATTACACATGGTGAATTTTACAAAAACATAATATGTGGATGATGTTTACATAAGTTTAT
ATCTTCTTCCATACTATGTAATGTGGTTCTACAAATGTTTAGGTAATTAGGGTTTAGGAGGGTATAATTAAATGATTTA
TTATTCAATAATATGCTTGTGTTGGGACATTGTGGAATTTTACCTGCTATTGTTGTGAGGCCCGGAGCCAAATTTAATC
TTATCTATTAGTGACAAATATATTTCTTAACCAGATTTTAAAGAAAATCTAGCCAAAGTTGTATGTGATTCATGTTGTA
TCTTCCTCCTGATGTTAGTTTAAATGCTTTAAGATAGTAGTAGCTATGATTGATGCCACTGAACTTTTAGTTCACTTG
ACTCCTCTATATAGCCCATGGATACCCAGTGTAGTACATTGATTTGGTTTTATGACTATTGACACCATTTTCTGATTTA
TGAAGGTTGACTTGTCCACAATCGGTGAGAGCTGGGATTTAATCCAGATATTCTGGCTACAATCCCAATAAGAGATGGG
TCATTCTTATCCTTCATTTCATGATTCATTTCATTTCATTCATTTCATTCTTTACACAGAAATGTGTTGGTAC
CTATGATGTACCAGGTATTTTCTAGCCATGGGAGGCATATCAACAAATGAAAATGATAAGAACCCCTACCCCTGTGGAG
CTTACATTCCAGTAGGGCAGAGGGGAAACAATGGCAGATAACATAATAAGTAAATTTCTGTAGTATATTAAGTGGTAGTA
ATGTATACTTGGAAAAAGTAAGCAGGTAAATGGAATTGGGCATACCAATTTTATACCTCTTTGGAGGGAGATATGAAT
TAATTTTTCAATTTTATCTCATAAATTATGTCAAAATAATAGGTTTTGTCCCAAGCCTTTCCAAGCAGGTAGCCTGGAA
CAAGTGTTCTGCTCTTCTCCTCTCTCCCCACTACTCAGAACATTGCTATAAAAGATAGCTAAATTACAAGATCAACTTA
CAGAGTCCTACTTAATTCATTATGTAGCTCAACTGTGGTTCAAATCTAGTAGTGTATAGACCTAACAGTCCTTACAG
TGGGTTTTCTCCCCAGTCTGGTAAACTGTATTCATGCTCCAGCTGCAGGTGACAGGAACCTCATCCTTTCATGCTGCT
CTTTTAGCTTTGGGAGTAAGCAACTCCCTCTCCTTCCACATTATCCAATATTGTGCGGCAGAGACTTGCTTCCATTA
GATACTGATAGTGGCTCCTCCACTGCTAGAAGCAGGAGGATGATCTTGGGGAATGATTATGGATTTAAAGGAGGAAGAG
ATAGTAGCATAGGCTTCTGTTTTACAGGAAATAGGAAGGTTGACAGTTGGAAGAAATCGTAGAGGAGTCCAGCTGGG
ATCAGTGACAGGAGGGAGGAAAAGGGAGGCCCTGGTCTCACAGGAAGGTTGAGTTATTGGGATGTTTATGAGTCAAGGA
TGGTTCGCTTATCACCCAGTATGTAGGTTTTCTGTAGATAAATTCCTGAGATGAATTTACCTATGCCTTCTTGCTTGATTT
CTTTTTTGCTTTTTCTTTCTTTCTCCCTTTCCCTTTCTTTTCCCTTTCCAGGGTATTGCTCTTTTGCCAGGCTGGAG
TGCAGTGTGCAATTATAGCTCACTGCAGACTCAAACCTCATAGGCTCAAGTGATCCTCTTGCTCAGTCTTCTGAGTAGC
TAGGACCACAAGCATGCACCACTATGCCTATCTAATTTTTTAATGTTTTTGTAGGGATGGGATTTTGCTATGTTGTCCA
GGCTGGTCTTAAGCTCCTGGCCTCAGGTGATCCTTCCATCTTGGCCTCCCAAAATGCTGAGATTATAGGTGTAAGCCAC
CATGCCTAGCCTCATTTAATTTTTCATAAAGTCTGAAATTATTATCTACTCTAAGTTTGGCAAGCAACTGGTTGTTTGA
GTTTAGTAATAATTTTTTGAAGATATAATGGATATAATTTTTTACATATTTGTTTTAATAGCATCCTCACAAAGAAATT
TTTAAATTTCTTTTATAGAATCTGATTATTTTACAGCCCTGAGGTACTCTTAATTTTAAATATATTTCTTTTTTAAAT
ACATTATTTTTTCATAAAGGCTTTATAATCAGCATGCTTTTTATTTTTTAAAAATATTGTACTACTAATATTGTTGCATAA
CTTCAAAAATTTATCATTTAGTAGCCATATAATACTGGCTAATAATTAACATAATATAATAATAATAATAATAATA
TAATATATAGTCTATTGATAAATAGGTATTTGAGTGTTTATCTGTTTTGCCCTTGTTTATACTAGGAAATGTGGCTTGCC
AAAATAAATGCATTATGCTTTCGGTTAGCTTGTAGTCTCATTCTATAAAGCAATGTAGATACAATGCAGTAAGGATAAGG
TGAAATCTGCCTTTGCAAAGGGATTTTTTAATTTGGAAAAAGAAGGGATTTGTGGGACAATGAGATAGAAGGCTTTACAG
AAAACTAGATGGCTTTGTGGAGAAAGAGAATTTAACTAGTTATTACCAGTTCAACCGCTACATCCCCATTACAGCCAT
CAGCATACTGTGTGCTTCTCTTTTCCCTGCTTTCCCTTGCAATCTATTTGCTATAACAGGTGCCACAGTAAAAAGCAA
TTCAGATCAAGTGACTCCTCTGTTTCAGAACCCACCAGTGGTTTTCTCAGGTCAATTTATAATGAACCCCTCAGTCTTTAAT
GATATAGCCTACAAAGTTCTGCATGATCTGCCCCCTTGGCTGCCTCTCTAACCTGATTTCTGACTTGTGTTTTCTTAC
ACACACTGTCTCTTGATGCTGGACGCATTCTGTACGTACACACTCCCAGCATGCTCTCACATTACAATGTTTGTAAT
TGCCATTCCCTCCACCTTGAAAGTTTGTCTCTGCAGTATTTCCCTGGCTTGCTCTCATTTCTTTGGATCTCTGCTCAC
ATGTTACCTCCTTAGAGAGGGCCTTCTGAAAATGATAGAAGTGTTTTTTTTTGGAGCACATCCCTCACTCCCCAGCTC
CTCTCCAGTGTATTTTTCTCCATAGCATCTTTCTGCCTGACCTTGATTTCTATATCTTTTTTGCTTTTTTATCTTCTGGA
AAACAAGCTCCATGAGATCACACAGGGATTTTGCTTTATTTCTCACTGCATGTCCCTCCTAATACAGTTCTTGGAATA
CAGGTGGGGGTTTCATAAATATTTGTTGGGTAAATGAAAGAATATATTAAATATATTATCATAAACCTTTTTTCTTTCCAT
GATTTGATGATGTGGAATTGAAACAGGTCTACATAGATGATAGTATATAATCAACAGCATTCCTGAGTACGTCTAGG
ATCAGCACTTGGTCTAGAAAAGATCTGTCTTCTGCCCTCCTTTCCCGTCCATGCACATCCACTGCCTTTGGGATCAAAC
CCAATCTCCTTCTCTGGCAGTCAGGGCCTTCTGAGACCTCATCTCTGTTGATCTTTCTAGCTTTACTTCTACCAAGTC
ATTCCCTCATAACTGCATGTCATACTGAGTGGCTTATAGTTCCCTCCTGTGGCCCTTGCAAGATATTTCTTGCAATTGC
TTTTCATTTGTCACTTATTTGTTTTATGGGTTTCTCCTCACAGTGCATAAGCTCTTCAAAGGAAGAGGCCTGGTATATCT
TTGTATTTCTGGGATCTAGCGCAGTTCTTGTATATTGTAGATGGTCAGTAAATACTTATTCAATGTAAAAGATGTTTTT
TGCTTCCAAAGACTAGTGGTAACCTGAAATGGTTGGATATAGGACAGATAAACTGATAGAAGCAGTTTTACCTCATTA
GAGGGCTTGATATAAGTATGTCCAGGGACAAATATAGTAAAAAAGAGAGCAATATTATAATATCCTTTAAAAA
AAAGAGACAGGTCTTGCTGTGTTGCCAGGCTGGTGCCATTATGGCTCACTGTAACCTCAAACCTCTGGGCTCAAACAG
TTCTCCTCCTTCAGCTTCTGAATAGCTAGGACTACTGGTGTGTGCCACCATGGCGGCTAATTAAATTTTTTTTTTTT
TTTTTTTTTTTTTTTTTTTTTGTAGAAATGGGTCTCACTATGTTGCCAGGCTGGTCTCAAATTCCTGGCCTCAAGTGATCC
TCCCATCTCAGCATCTCAAAGTGCTGAGATTATAGGCGTGAGCCACTGTGCCTGGCTAGAATATCCTAACTTTTAGCAA
AATCATACCTGAGATGAGGAAGGATGGCATATAACATGATTGAATCAGCACACTTAACTAAGAAAGATAGCCTAAGTGG
AAAGAAGGGTTATTTGAAAGGCATTTATTTAGAAAGAAGGGGAGCCATTTGGCAGTGGACTGACTATGTGGGCATCAGTC
CTTTATTTAGTCACTGAGAGTCATGCAAGGTGGTGAGCCTGCTGATAATGAGGAAATAATACAGAGGGGAGAATTCAGG

Fig. 9.5

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GGAGTGATGACACGTTTCACGTTCTTGCTATTTTGGACTTATTAACAACGTGAGTGTTGAGGGTTATGATAGTGTTAGCC
TGGCTGCTCCTGGTTCGGGAAAGAGTGCTATGAAAATTGAGATATATCTGTCTAGTGATGGGAGGAGAAAAATGGCTGT
ATTTGGTTCCTATATATTTGTTCATGCCTGGTTGATGGTTTGTATTTTCTATAGTGATTTGTGAGTGACAATGGATCTATA
TTCAATTGACCCAGTATAGTTGAATATGTGAAAATATTTTATGTAAATTTTACTAATATATATTTTGTAAATAACAACC
ATATAGAATCTACTAGGCTATAGGTGTTGAAAGATAACATTTTTTTTTTTCTAAATCAGGTTTGTACAGAAGGAGAAA
GTGGCAGTTGAAGCATTTCAGATTGCTGCCTTCTCCTACCTCCTGAAAATAGGAGAAAGTTACAGCTATTGATGAGGA
TGATGGCAAGGATTGCTTAAACAAAGAGATGCCACCCCTGTGTGATGGCTTTGGTACCCGAACACTGGTAGGTTGATT
TTTAACATCAGGTATGACTTTTTGGAATGAAAGTCGACTGAGTAAGGTGTATTAGTTTGGGTGATCAGAGGGAAGTACA
AAGCAGACTTGCTTAAAAGAGCCACAGTCAGTGTCACCCAGACTTGTGTCGTGCTTTTGTCTAAGTGCTTTAAAGAA
CATAATGAGAAATGGAGACTTGAGTAGGGGCAGACTCTGTAGAGAGATCTGCTACATCCTGTTCTCCCACTTAAATTA
TTTTAAATTTATTTCTTAATTGACAACATAATAATTGCATATGTGATGTTTTAATTCATGTATACATTATAGAGTCAATCA
AGCTAATTAACAGATTCATTACCATAACCACTTATCATTTTTTTGTAAATGAGATCATTAAAAATCTATTCTTTCTCAA
TTTTGAAATATACAATATATTATTATAATTGTCGTACCGTGCTGTGCACTAGAGCATTAAGACTTGATCCTCCTGTCT
CACTGGATCTTTGTACCCCTTTGACCAACATCTATTTCCATGTCCCTGTATCCCCCAGCCTGTGGTACCTACCATCTAAT
CTCTGCTTCTATTGGCATATGTTTTTAAAGAGAGTTTATTTTAGAATTTACTTACTAGCAAAATCTCTTGATTTTCTC
AAAAAATGTCTTTGAGCAAAACTGGTGCTTCAGTGGAATGTATGTGCTTTCCATATCCTGTAAACACCCGTGTACTCC
CTGGGGATTCACTGGCCCAGTTTTAGAAAGTAATAGAGTGAGATCAGAGGAGCACTTAGTGCCCAGTAGGATTCTTGATC
TCTATATTTTTTTGGAGTGAGTGGATGGTTAGAGAGGCTAATAGACCTTTTTGAAAAGCTGAAGAAAACCTTTATATCTA
TTCTGCATATAAAATACACATGCAAACTTTGCATGCAGTTTCAAGGCGATCACAGAGTCCCTGGGGTCCATGGTCTCC
AGGCAAAAAATCTCTGCCTGTGTGTCTCTGTACATGTAAATGTAAGTGAAGATCATTTCAGTGTTAATGAAGT
GTTACTAAGCAGAATTACATGATTTTTTGTCTTAAAGATTTTGAAGGAAGATTGTCAGCAATAAAAGTAGTAGGAAAAA
CAAAACAAAACAAAAGCCAGTCCAAAGAACAGTATTAAGAATGGAGTAGTTAGGCTGGGCGTGGTGGCTCACTCCTGTA
ATCCCAGCACTTTGGGAGGCCGAGGCAGGTGGATCACCTGAGGTCAGGAGTTCTAGACCAGCCTGACCAACATGGTGAA
ACCCCGTCTCTATTAAAATTACAAAATTAGCTGGGCATGGTGGCAGGCGCCTGTAATCCAGCTACTCGGGAGGCTGG
GGCAGGAGAATTGCTTGAACCTGGGAGGTGGAGGTTACAGTGAGCCAAGATCACACCATTGCACTCCAGCATAGGTTAC
AGAGCGAGACTCCGTAAAAAAG
ATAATATATTTTGCATTTCTGCTTGCCCTCAAAGAGAGTCTCATCACTTAATGTACTTCTTGAAAAGTCCCATGGCTTTG
TTCAAAGATGATTGTCAAATATACATCGAAAATCCTAGAGAAAATAAGCTAACTAGGAAAAGATGTAATGGTTGAGGCA
TGTGGACTGTGTTAGCGGTTTAAACAAATCAATAATGAAATTAGGATAAATAATAATTGACTAAATGGCTATAAAGTG
CTTTGAACTTAGTAGTGCTACATGATTCAAGTCATGGTGATAATGATTTTCAAATTTGTTTTCATATATTTATATGTG
TGTATGTGTGTAT
ATTGCTCAGATAAGCATCTCTCTGCAGTCTCTTAACAGGAAGATGCACATTTTGTGTTGTTCTGGTTGTTGAGACCGTCA
ACCCATCTGAGATGGCTGGGTACTCTTGGTAGAATATGCACCCTGTGCCTTAGACCATTCTCATATTTGCCTTTTGCA
AATTAGATTGCACTCAGTTCAGAAAAATACTCATCTGCGTTTACCCCCCACCCTGCTTTTTTTTAGATGGTTTCAACA
TTTTCCCGTTGCATCTTGTGTTCCAAGGATGAAGTGGACTTGATGAGTTATTAGCTGCTAGATTGGTAACGTTTCTGA
TGGACAATTACCAGGAAATTCTGAAAGTCCCTTTGGCCTTGCAAGCCTCTATAGAGGAGCGTGTGGCTCATCTACGAAG
AGTCCAGGTAAAGGAGAGAGATATTATCAGTTCTATGAAATTGGCAATATAAAGTCACGTAAGCTTGCTAGGCTTCTTGG
GGCATTTTTATTATGAGGATTAAGTGAATTAATGTTTATAAAGTACTTACAACAGGTCTTGGCACATAGTAATCCCCTGC
ATGTGTTTGTATTATCATTGAAAAGTTTACAGAATGCACATTTGGTTTTGTGTATCAATCCATGTGGCATATTTTTTAT
TGCTAATTCCTTGAATGAGTTTGTTTAGAATCTACATACACCTACTGATGGAAATGATTTTATGTGCAGTAAGTGCTT
GATTCATAATTGCCCTGTGACCTAGTTTAATATTCCCCTTTTTGTCATGAAAAGCTAAGGCACAAAGAACTTAAATAA
CTTCGCTTAAGGCCACAGAGCCAGAATTCTGACTGTTGCAGTCAGCCTCTGTAATCAGTGCTCTAAACTATCATTCCAT
ATTGCCTGTCTAAAAATCATAGTACAGTAGAAGAACACTGGCATGAGAGCCATGAATCCCTGGCCAAATTCCTAGCTGT
ATCCTTTCTGTGCTGTGACTCTGGGCAAGTCCCTTAACCGCTTTGGATTTTACTTTCTTTCTCCTCAGGCGTTAAATGGG
CTATTCTTTCTTTATAAAGTTGTGAAGATTAGAACACTGTACTTTACAATTCCTAGGATGAATTTCTGACATATGTCAG
GTATTCTTTGAAAGGTAGCTGCTGCTGTTGTGTCATGGTGGTTATATAAAACAATGCAAAAGAAATATAATATATTATTA
TCGTGTGGACCCCAAAGTGGGGTTTGTATGTCAGCCTGTGGGAGCAGAAACCTGGTAAAGGCAGGCAGAGGTTCACTT
GACTCCTTCTGACTGATTGGACCTCTTCTCATTTGAGATCTTTTTGAGGCACCACAATCAAGGACCAGTGAAGTGCCTC
CTTCTGAGGTATTACAAAATAATTAATTGTCCAGCTGTTTCTAGAAGGCAATTTTAAAATTTAAATTTTTTCTATTTT
TTTGCAATTATCAACCTAACACATTGAAGAACTTGGGAAATATAGAAAAACACACCAAAAAGTATCAGAGTTACACT
AAAATGTTCATAGTCAGTAATATAGTCTTTTCTAGATTTTTTTTTCTGTGTCTGTCTGCTGTCTATGTCCATGCCTACCTT
ACTTTCTTCCCTCTCTCTTCCCTCCCCACCCTCCCTTCCCTTCCCTCCTTCCCGCCTTCCCTGCTTCCCTTCTTGCC
TGCTTCCCTTCCCTGCTGCTGCTTCCCTTCCCTGCCACTTACATTCTGTAAACAATGAGAAAAAATTTCCCCCATCATTAAT
ATTGTTGATTGGCATGAGAATGGCCAAGAGCACAGACTCAGGAGCCTTTGAATTCATGACATGCCTTTTGTAGCTGTG
TCACCTTAGGCAAATTATATAACATGTCTGTGCTTCAACATCTGTCCCAACATTTCTTTTTCTCATCTGTAAAATGGGA
ATAATAGTAGTTAATATTTACCTCATAGAGTTTTTCTGGGGAATTAAACATGTGATTAAATAACATGTGGGCTTAATA
CATGTGAAATGCTCACAATAATGTTTATCACATTGTAAACCTACAATTAGTAGCTGCCTTTGTGTTGGTATCATCATT
ATTGTTATATTATTCATAATAATATTATTGGCAACATAGTATTCATAATATGGATATACCATAATTTATTTAAATGA
TACTTTTTTGGTTGGTATTTAGATTGTTTTTCATGCTTTCCCCCATTTTGGGAAGCAATATAGCAGTAAATATTTTAAAGG
TAGATTTTTTTTTGGCTAATCTGTGATTATTTTTTAAGAATAAACTCCTAGGGGCAGCATTTGCTTGGCCAAAGGCCATGA
ACATATTTTAAGTATCTATAGCATATTGCCAAATTTAGAATGATCATTTCAATTTACATTTCTGTGCTAGTGTTATAAGAG

Fig. 9.6

AGTGTTCATTTCTTTGCCCCCTTTGCCTACTTTGGATATTATCATTTAACACTTGTATATATCTTTGCCACTTGCATGGGT
GAAAAGTGTAATTTTAACTGCTGTTTTAGTTTTAATTTTTTCTCTGATATTTTTATGAGCCAACCCTAAAGAAAATAAA
AATGAACAGAAATACTTCACCAAGTTTCTGCAAGGAAATGTTATAGCAGTGTATTAGTCTGCTCTCATGCTGCTAATAA
AGACATACCCAAGACTGGGTAATTTGTAAAGGAAAGAGGTTTCATTGACTCAGAGTTCAGCATGGCTGGGGAGGCCCA
GGAAACTTACAATCATGGCAGAAGGGGAAGCAAACACATCCTTCATGAGGCAGCAGGAGAGAGAAGCACCGAGCAAAG
GGAGAAAAGCCCCTCATAAAACCATCAGATCTCATGAGAACTCACTCACTATCATAAGAACAGCATGGGGGGAACCACC
CTGTGATTCAAGTACCTCCCATCTCCACAGCAAGGGGATTATGGGAACTACATTTCAAGATGAGATTTGGGTGAGGAC
ACAGCCAAACAATATCAAGCAATAACTGTGTGTCCTGTATACTTTGCTAGGTTTGTGTATTACATGCATAGCACAGGC
ATATATTGTAGTAACTTAGCCTTGTGAGCCCTTTGCTATTACTTGAAGTTCAGAAGGCTGAGCTATGGTGATTAAATTA
ACTGCAGGTAAACATGATCTTGTTAAGAGACACTGCAGTGTGCTCTGAATAAAATCAGTAGTGATTCAATTTGTCCAGTT
ACCTTTTCTCTCTTGCAAGTACATATTAGAATTGCCAAGCACCTGTTCCATTACAGCCCTCAGAACCTATAGATCTTTGT
TCTTTTGATTAGGCTCCTAAGCCTCTACACTGTATCACATTTAGGGGAGTGCTTCTCTGAAAATGCAGTGTGCTTGGA
TTATTCCAGTGATTGTTTTAGGACAGTTTCAGGGCTGACCCTGGGATACATTATTAGGTAGGCAATGTTTTTGACACAA
GCCTTCTATGATCCTCATCTGCAACCTTCTATTTCAAAGGACTCCAGGCTCTGAAATTAATAATTTTTTAATGTATGTTT
ATTTATTCACAGGCCTTGGTTTTCTTTTTCTCTTATTTCTCTCTTCTCCCTTATCCCTTCTGCAATGTCTGCACCTTT
GTTGTTTTACTTGATTCTCATGGGTAAAGATGTTAGGAAATGATTGTAGTACCCCTCTCTTCTTAAAGCTTAGCTAAAT
GCNTGCCACTACCATCCCCAAGGCATTGAGAATCACACTCTTCAGATGTGGGAATGCGCCTGGATAGTTCCAGTGGATA
TCCAACATTATCAATGTTTTGAATCANTTTAGTCAATGTGTTTAAATTTATGCTTGAATTTAGATCTTGTTAAAGAGG
GAATGCCGAGACCAGCTCGGTCTGGAGACCCTAACCAGTGCCCTAGAGGAATTAAGACACACACACACACACACACA
CAAATATAGAATGTGGAGTGGGAAATCAGGGGTCTTACAGCCTTCAGAGCTGAGAGCTTTGAACAGAGATTTACCCACA
TATTTATTGACAGCAAGCCAGTCATAAGATTTACTAAAAGTATTCCTTATGGGAAATAAAGGGGTGGGTGAAATAAAGG
GAAGAGCTCTGGCTAGTTATCTGCAGCATGAACATGTCTTTAAGGCACAGATCGCTCATGCTATNGTTTGTGGTTTTAAG
AATGCCCTTAAGCGGTTTTCCGCCCTGGGTGGGCCAGGTGTTCTTGCCCTCATTCCTGTAAACGGACAACCTTCCAGCA
TGGGCATCAAGGCCATCACGAGCATGTCACAGTGCTGCAGAGATTTNGTTTATGGCCAGTTTTTGGGGCCTGTTCCCAAC
GAGGGCACAAGCTTTTTTTTGTATTGAAATGGCTCTTGGGTTTACAAAGGTAAAATCCAATTTAAACCGTGGCTTTA
AATAATAATATTTTGAAGTTATGGAAATACTTTTACTTTTCAATATAGAGTTTCCCCCAATCAAATTTTGAGGACTAAC
ATGGCATTGCTACGGTACAAGGAGATATCCTGTATTTGATCTAAACTAATTCTTATAATTTATTTGAAAATATGCTATA
TTCTTTCACTACTAATCCTTTTGCCCTCAGATAAAAATACCCAGGAGCTGATATGGATATCACTTTATCTGCTCCATCAT
TTGCCGTCAAATTAGTCCAGAGGAATTTGAATATCAAAGATCATATGGCTCTCAGGAACCTCTGGCAGCCTTGTTGGAG
GAAGTCATAACAGATGCCAAACTCTCCAACAAAGAGAAAAAGAACTGAAGCAGGTAAAAGGATATTTCTTTATAAG
CCATTTTTTCTCCTCCCTACTGTTTTTTATGTTTTTTTTTTTTTATTTAATAGGCTTCATGTAAATCTCATCTTCTGAGCC
ATAGCAGTTTTCTAAGTATATTAGTTCATACATTCTAGAAGTCTCACTGATTTACACACTTGTGGTCAGTATCCTAGAT
CTTTATTTCTTCAGTGTAATCTGTTACCATTGCCAAACTCTTCATTTTTTGAGCTATTCTCTAATATTTGTCCTTTGTT
TTGGTATGTGGCATGATGTGAATTAGCAGAGCATCTGCTTTAATCTCAGAGAAGCTATCAGTGAGGGTGGATTTCTTTT
CATGTTTTCAAAGGCCAAATCAGATGGAATTCACAGGTGTAAATGTTTAGAGTTTAAAGCCGTTGGTTTTAGATAGC
CACTATCAAATTAGTTTCTTGCCCTCAGGTATCAGTACAAGTTTTTAGTTTCTCTTATGTTAGGGGAACACAGCTTAGGG
TTCCAGCGAATCGTTTTTAACTCACAATATGTTACTTTTGACTATTTCATAAAGGTTGATCCATGATCAGGTGAGTGGTT
TTTGTTTTTAAAGACTGAACTCTACATAGTTGGATTGAGGGAACACCGTACATGTGGGATTACACACAGAGCACCAGAG
TTCTGGGCTTCTGATCATATTATTCTCAGAGTAATTTTGGAGGAGAGTTTAGTGGTAAAGATCAACAGAAAGTATACTG
CCTCTCTTTTAAAGCTTTTCACAAAGTAATATACACCAGAATATTCCTCTTCCATTTGAGCAGAAGCTGGTGGGGCTGA
GTTGGAGAAACAGAGAAATGATGAAATGTAATTGGGCCAAAATTTTATTTAACTGTGTAGAACAGAAGTTATTTTGAAG
TCAGAGTTACTTGTGAATATATAAAAAGTGAGGTCTATTTATAGTTGGCAGTGGGGGGAATTTTCAGAAACATGATTAA
AAGGGGAACCTTATTTGAAGCTTAGATCAAAGGAAAGATGGATTAGCTCTATAGAAATTGGAATGAAATGTTTTAACAAA
AATAAACAAAAACAGGAAAATACAGGCTCAGAACTCTTAAAATTTGGCTGCCAAATTTTATTGGCTTTTATGTGTCTT
TTTTTACTCTTCTTGGACCATTCATTTTTTTTAGTTTCAGAAATCCTATCCTGAAGTCTATCAAGAACGATTTCTTACAC
CAGAAAGTGCAGCACTTCTGTTTCTGAAAACCCAAACCGAAACCACAGCTGCTAATGTGGGCACTAAAGAAGCCTTT
CCAACCATTTCAAAGAACTAGAAGTTTTCGAATGTAATAATACTTCCACAGCAACAGGTGCTAGAGACCACTGTTGTTG
TTTTGAGTGAATGGTGGTTAGGAGAAAGACTTTGGTGGTGGGAGAAAGAAAAGCATAAAACAAAGACTACTGAAATATA
GATAAAGATTGCCTTAGTTTTTAAAGTGTGTTGGCCATTAGTATTTTTTATAAACTCAATGCTAGTTTTTAAAGTGTATA
AATTGGTTAAATTTATGAGTCAAATATATAGTGATAATGTTAACATGTTTGTAATTGCTACAGAATTTAAGGGTATTT
TTATCTCTGTGCTTTCTTTTTTCATGGTGTTTATTAAATAATTGTGTATATACATCCTAGCTACTGATATCTTTATTATA
GCCTTAAGACTTAATTTTAAAGTCTTAAAAATAGCGTGTATACTTGAATAAGAAAGACACTGGGTACTGTACTGTGATG
CTATTGACTTAGTAGCCAATTATCATTTCTCCTGTATAAATTCCAGTTTTTTATTGCTGCACATAAATTTTTTAAATGTCT
TATATTGTGATAGCTATGTCTTTTATTGCAGATTTATTGGATGTTATGACAGATTTTACTAAAGCTAGTGTTTTTATAA
CATATATATTAGTTGATGTTTACCTATAAGTGGAGTAGATTTTCATCTGCCTGCAATGGTATAATTTCAAGTCTTAGCTA
AAAATGGAAAGTTGAACTGGATAAATTTCTTTGGGTACCCCTTAGACCTCTGATTCTAAGTCAAATGCAAATGGGTAAAT
AAAATGAGACTACTTCCCTTTATAAATATATTTTCATCCTTTTGAAAGTAAGTGAAATGTAATAAACTTATTTTTTTTA
AAAATGCCATACCTCGTGTCTTTTTTTGGCATTATTTTAGATCAGTTAGAGAGCATCCTCAAATGCCTTTGCCAATGG
TCTTCATCAGTCATTTATTTCTATGAGAAAATGAAAAGTAAGATTTGGGCTTTTGTACTGACTGGTCTTTAGGTGAGCA
CTTTGCCAATTTTTTTAGCATTGCTTTGTCTTTTGTAATAATTGTGGTGCCTTTGGATTATGGAGCATCTGAAAGTCTTC
CTCTGAGCGCAATTACCCAGAAAGCTGAGAGCAACACCCACTAATTGGCTTACAGTACTCCTTGGACTGATTTTTCTCT

Fig. 9.7

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GCACTTAAGATTTGCTCCTGTTCTCTTGTAGCAAAATACCTCCTACATCTGATGCAGATTTTGCTTTTAAAAATGGACC
AAAGTATTTCCTATTGGTTTGGGTACCACCTTACATTCCAAATACATAATGTAATGGGAGATTTTGTAGTGTTCAGGATT
CATTCTCAGACTTTGGCATTCTGTTTAGAGCCAAGAATAATTCTTCTTTCTGCTCTATTTTCACTCCGAGGTAGAGTTT
ATTTTTCAGAATCTCTTTTCTGTTGCTTGTCTTTGGAAGTATATGTGGTTGTTTTCTTTCCTTTTTTTTTTAAGATTA
CTTTTTTAAAAAATAACATATTTATGGAGAGATCATAAACATAAGCACCAATTAAGTGTGAAATTGATGAATAGA
TAATAGCAGTTATCCAAAGTGGTTGTCCCAATTAACGCTCCCATTAACAGTGTATGAGAACTCCTTTTGTTTAATATCC
TCACCGAAACTTGCTATCGTCAGAGTTTGAAATGTTGTCAATGTGGTGGTTATTAATAGTGTCTGATTATGGTTTTATT
TTGCATTTCCATGATTATTAATATGGCTGAACATTTTTCCTTTTTCCTTTTTCGCTCTTGTACAAATTTTGGT
ATCAGATTGTCAAGCCTNCATACACACACACACATGCAGAGACACACACTCACCTGTGAGTTTTTTTTTTTAATT
TTGATTGTATTATATGTATAAATCAATTTGCAGAGAAATGCTGTCTACAATATTAGGTCTTCTAGTCCATGACCATGGT
TTACCTTTCCATTTTATTTGAAATAAATGAGAAAAAAGTTGCTTTTTTCTGTGAAGATTTACAAATCTTNGTTAGAT
TTGTTCCCAAGTATTTGATACTTTAAAAAAGTATCAAAATTTTCAAAGACTGTAGTACCCATTTCTGTTCACACAGT
TACTCAGCTGATCAATGTTGACGTTTAATCTTTTGTAAATGCTGGTGTGTAAATTTCTTATAGTGTGTGTGCTTC
TGTATGGTATAAACTGATTTTTCTGTAGTTTTGTTTACAGTGAACCTTGCTAGACTTGTTAATTCTAAACATTTAGTA
TGATTCTTTGGCATTGTCTACNTATTAGGAAAAAACAGCTTTTGCTTTTCCGTTCCATGGTAGACAGCATCTATATTG
TCCTCAGTGATCCTTGCCCTCCTGGTATTTAATGCCTTGTGTAATTTCTGTCTTGAGTGTGGGTGGAGCTAAAATGAC
ACATTTCTAAATAATAGAATATATCATCAGTGATGGGGTATCACTTCTCAGATTAGATTCCAAACCTCTGGCTTCCATC
TTGCTTGTCTTCTTCTTATTTCTCTCCCTGGCTTGCTCTGAGGAAAGCCAGCTGTCATGTTGTGAGCTGCCCTTTCAGA
GGCCACCAGGCAAGGAAGTGTCTCTGGCCAACAGCCAGCGTGGACATGAAGCCTGCCAACAGCTTTACGAGTGAT
CTTGGAAGTGGATCTTCCCTTGGTCAAACCTTGAGGTGACTCTACCCTAGCTGACACCTTGATTGGAGACTTTTGAGAA
ACTCTAAGCCAGAGGACTTAGCAAAGCTGTGCCTGGATTCTTATTCACAGAACTGTGATATAATAAATGTTTATTGT
TTTAACCCACTTAGTTGAAGAATAACTTGTTATAAAGTGACCTAGATACAATATACTTTCCAATCTCCATACCCCTCCC
TCCCTTCCCTTCCGTCCTTCCCTCCCTTCTCTCCCTTCCCTTCCCTTCCCTTCCCTTCCCTTCCCTTCCCTTCCCTT
CACTTTCTTCTTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCT
TCTCAGGACACTGGAGATCTTCACTACAGCAGTTCCCTGGTTCCCATTTATCTGCTGTTTCACTTTGTACAGTTTCAGT
AAGCCACGCTCAACCACAGTCAGAAAATATTAAATGGAAATATTCCAGAAATAACAAGTTTAAATNGTGTGCTGTTC
TTATAGCATGATGAAATCTTGAGCTGTCTGCTTTGTCCACTCTGAAAATGACTTATCCCTTTGTTTCATGGATCCATG
CCTGGTAGACCAGGTACATTGTNAATGAGCAGTAATATTTTGAAAGGAATCTTTTTTTTTTTTTTTTTTTTTTTTGAGCA
GCAGATCTCAATAGTGGGCTTAAATATTGAGTAACTATGCTGTAAACAGATGTACTGTTATCCAGGCTTTGCAGAGC
ACAGGCAGAGTAGATATAGTGATTTTTTTAAGGGCCCAAGGATTTTTTGAATGATAGATGAGCATTAGCTTCAGCTTCA
GGTCACCAGCTGCTAACAGGAGAGTCAGCTTGTCTTTGAAGTTTGAAGCCAGGCATTGGCTTCTTTATAGCTATGAA
AGTCTTAGATGACATCTTCTGCCAATAGAACGCTGTTTTATCTACAGTGAAAAGCTATTGTTTAGTGCAACCACCTTCA
TCATTGATCTTAGCTAGATTTTCTGGATAACTTGCTGCAGCTTCTCCAGCCTTTGTTGCTTCACTTGCAGTTTTCAGT
TAATGGAGATGGCTTCTTCTTAAACCTCATGAACCAGGCTTTGCTTGCTTTCAACTTTTCTTCTGCAGTTTCTCAC
CTCTCTCAGTCTTTATAGAATTGAGGAGTTAGGGCCTTGCTCTATAGTAGGCTTTGGCTTAAGGGAATATTGTGCTGG
TTTTATCTTCTGTCTGACCACTCAAATGTTCTCCATTTCAAGCAATAAGGCTGCTTAGCTTTTGTATCGTGTGTGTTA
CTGGAGTGGTGCTTTTAATTTCTTCAAGAACTATTCTTTTCACTCACAACTTGACTAACTGTTTGGCACAAGAGACC
TAAACTTTCTTCTATCTTGGCTTTCAACATACCTTTCTCATTAAGCTCAATCATTTATAGCTTTTCGATTTAAGTGAG
AGGCGTGCAACTCTGTTTCACTTGAACAGAGGCCATTGTAGGGTTATTAATTGGCCTGACTTCAATATTGTTGTGTCTC
TGGGAATAGTGAAGCCCAAGGAGAGGGAGAGTGATGGGGGAATGGCTGGTGGGTAGAGCAGTCAGAACACACAACATTT
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TGTAATAATATGCAGTGATCTGGATACAATATAACAATATAATCTGTGAAGTGCAATGAACCAGATATGCCTGTGTATAT
AGTGTTCAATACTATATGGTTTCAGGGCCAGGCATGGTGGCTCATGCCTGTAATCCTAACACTTTGGGAGGCCAAGGCA
GGCAGACTGCTTGAAGCCAGGAGTTTGAGACCAGCCTGGCCAACCTGGTGAAATCTGTCACTACTAAAAATTTAAAAA
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GCAGAGGTTGCAGTGAGCCCAGATCACACCACTGCACTCCAGCCTGGGTGACAGAGTGAGACCCTGTATCAACAAAAACA
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GGAATAGAGTGGGGATCGCAGCAGCATCATCTTGCTCCTAATTTCAAAGAGGAGGTTTTTAGCATTAGAGTATTTGTAG
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GGGGTTATATTACATTAATTGATTACTAATATTAAGCCACTTTGCATTCTAGGAGTGGCATAAATCTAATTATGATGTA
TTATCATTTGAATATATATATCTAGATTCTGTTGCTAATATTTAGTTTATAGTTCTTGTATCTATGTTTATGAGTAAAA
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AATTACACCGAAGTCAGAAAAATGGTTTGTATAATTTCAAGTCTTTGAAATTTGTTGAGACTTGTTGCATAGCCTAGTA
TATGGTCAATTTTTTGTAAATTTCTGTACACATGAAAAGAATATGAATTTTGCATTGATTACAGTTTTTTTTCTCTCTCT
ATAAAATCTTGTTTAAATTTCTTACAATTTTACTGATTTATTTTGCCTGCATATATATGTGTGTGTGTGTGTGTGTGTG
TG
CTGGAGTGCAGTGGTGCAGTCATAGCTCACAGCAGCCTTGGCCTCCTGGGCTCAAGTGATGCTCCTGTCTCAGCCTCCC

Fig. 9.8

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AAGTAGCTAGGACTAATTATGCTCAGCTAATTTTATTCAAATTTTTTAGAGATTTTGTCTTGCTGTGCTTGTTACCCAG
ACTGGTCCCAAACCTCCTAGTCCTAAGTGATCTTCCCACCTTGGCTGCATTTATATTTATTATTGAGAGAATTATTATAG
ATTTCTGATTAAATTCATTTTGAATTTTCATATTTATCTGGTGGACTGACCTTTTTATTATTATTATAAAATGTCATT
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GGCTTCCATTACTGCTGTTTAAAGTCAGCTGCCACTCTAGATATCACTTCTTTATCAATAATGTGCCCTTTCTTGGC
AGGCTGCTTTTCTCTCTGTCTTTCTGTTGCTTTTCTTCTGTTGCTATATCTTGGTGTGTACTTTTTTAAATAATTATCC
TGCTTGGGATTTCTTGAGCTTCTTGAATCAGAAGTTTGAGGTCTTTTGTGAGTTTAAATATTGCCTTTGTCAATCTC
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GGTCTGACAACAGACATTTTCAAGTGATAATTTACCAAGATAGCAGAACTGTTTGTGCTAAAAATAGTTATGCAGGATA
AAATGCAATCTATAATTTTATGAGAACCTGTGATTCTAGTTTACCTCCAATTACTAATATCCATATCTATGTTTGTATA
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Fig. 9.9

TTGCCTACTGATTCTGCATAATTTTAAAAATTAAATTTGGGTTTTTGTCCATATGCAAATTTGTCCTAAAAATTTCCAT
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AAGTTTTTCAAATGACTCATGTGATGCCTGGTTTCAAGCTTTATCATTAAGTCAAGACCAAAGTTATATGGGTTTTT
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ATAACATGCTGGTCAGGTTGCAGAGAAAAGGGAATGCTTATACAACTGCTGGTGGGAGGTAAATTAGTTCAGCCACAGT
GGAAAGCAGTTTGGCTATTTCTCAAAGAACTTAAACAGAACTACCATTCAATCCAGCAATCCCATTAAGTGGTATATG

Fig. 9.10

[illegible]

Fig. 9.11

CTGCTTATCATTTTGAATTCAGCTGGTCTTGCTAAAGCACATACACATGGCTCTTCCATGTGACCTCTGTGTTTGT
GGGCTTCCTTACAGCAGGGTGGGTGCCTTACAAGAATAAACATCACTGAGAAAACCAGAAGAAAGCTGTATTGCCATTT
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CTACTGGGTACATAACGAAATGAAAGCAGAAATAAAGATGTTATTTGATACCAATGAGAACAAAGACACAACATACCAG
AATCTCTGTGACACATTCAAAGCAGTGTGTAGAGGGAAATTTATAGCACTAAATGCCCACAAGAGAAAGCAGGAAAGAT
CCAAAATTTACACCCTAACATCACAATTAAGAAGACTAGAAAAGCAAGAGCAAACACATTCAAAAGCTAGCAGAAGGCA
AGAAATAACTAAGATCAGAGCAGACCTGAAGGAAATAGAGACACAAAAAACCTTCAAAAAATTAATGAATCCAGGAGC
TGGTTTTTTTGAAGAGATCAACAAAATTGATAGACCGCTAGCAAGACTAATAAAGAAGAAAGAGAGAAGTATCAAATAG
ATGCAATAAAAAAATGATAAAGGGGATATCACCACCAATCCCACATAAATACAAATTACCATCAGAGGATACTACAAACA
CCTCTACACAAATAAACTAGAAAATCTAGAAGAAATGGATAAATTCCTGGACACATACACCCTCCCAAGACTAAACCAG
GAAGAAGTTGACTCTCTGAATAGACCAATAACAGGATCTGAAATTGTGGCGATAATCAATAGCTTACCAACCAAAAAA
GTCCAGGACCAGATGAATTCACAGCCGAATTCACCAGAGGTACAAGGAGGAAGTGGTACCATTCTTCTGAAACTATT

Fig. 9.12

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CCAATCAATAGAAAAAGAGGGAATCCTCCCTAACTCATTTGATGAGGCCAGCATCATCCTGATACCAAAGCCTGGCAGA
GACACAACCAAAAAGAGAATTTTAGACCAATATCCTTGATGAACATTGATGCAAAAATCCTCAATAAAGTACTGGCAAA
CCAAATCCAGCAGCACATCAAAAAGCTTATCCACCATGATCAAGTGGGCTTCATCCCTGGGATGCAAGGCTGGTTCAAT
ATATGCAAATCAATAAATGTAATCCAGCATATAAACAGAACCAAGACAAAAACCACATGATTATCTCAATAGATGCAG
AAAAGGCCTTTGATAGAATTCAACAACCCCTTCATGCTAAAAACTCTCAATAAATTAGGTATTGATGGGACGTATCTCAA
AATAATAAGAGCTATCTATGACAAACCCACAGCCAATATCATACTGAATGGGCAAAAACCTGGAAGCATTCCCTTTGAAA
ACTGGCACAAGACAGGGATGCCCTCTCTCACCCTCCTAGTCAACATAGTGTGGAAGTTCTGGCCAGGGCAATTAGGC
AGGAGAAGGAAATAAAGGGTATTCAATTAGGAAAAGAGGAAGTCAAATTGTCCCTGTTTGCAGATGACATGATTGTATA
TCTAGAAAACCCCATNGTCTCAGCCCAAAATCTCCTTAAGCTCATAAGCAACTTCAGCAAAGTCTCAGGATACAAAATC
AATGTACAAAAGTCACAAGCATTCTTATACACAAATAACAGACAAACAGAGAGCCAAATCATGAGTGAACCTCCCATTC
CAATTGCTTCAAAGAGAATAAAAATACCTAGGAATCCAACCTTACAAGGGACGTGAAGGACCTCTTCAAGGAGAACTATAA
ACCNNTGCTCAATGAAATAAAAGAGGATACAAACAAATGGAGGAACATTCATGCTCATGGGTAGGAAGAATCAATATC
ATGAAAATGGCTATACTGCCCAAGGTAATTTATAGATTCAGTGCCATCCCCATCAAGCTACCAATGACTTTCTTCACAG
AATTGGAAAAACTACTTTAAAGTTCATGTGGAACCAAAAAGAGCCCGCATTGCCAAGTCAATCCTAAGCCAAAAGAA
CAAAGCTGGAGGCATCACGGTACCTGACTTCAAACCTATACTACAAGGCTACAGTAACCAAAACAGCATGGTACTGGTAC
CAAAACAGAGATATAGATCAATGGAACAGAACAGAGCCCTCAGAAATAATGCCACATATCTACAAGTATCTGATCTTTG
ACAAACCTGAGAAAAACAAGCAATGGGGAAAGGATTCCTTATTTAATAAATGGTGCTGGGAAAACCTGGCTAACCATATG
TAGAAAGCTGAAACTGGATCCCTTCCTTACAGCTTATACAAAAATTAATTCAAGATGGATTAAAGACTTAAATGCTAGA
CCTAAAACCATAAAAATCCTAGAAGAAAACCTAGGCATTACCATTCAAGGACATAGGCATGGGCAAGGACTTCATGTCTA
AAACACCAAAAGCAATGGCAACAAAAGCCAAAATTGACAAATGGGATCTAATTAAACTAAAGAGCTTCTGCACAGCAAA
AGAACTACCATCAGAGTCAACAGGCAACCTACAAAATGGGAGAAAATTTTGTCAACCTACTCATCTGACAAAGGGCTA
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GACAAATGGGATCTAATTAACTAAAGAGCTTCTGCACAGCAAAAGAACTACCATCAGAGTCAACAGGCAACCTACAA
AATGGGAGAAAATTTTGTCAACCTACTCATCTGACAAAGGGCTAATATCCAGAATCTACAATGAACTCAAACAAATTTA
CAAGAAAAAAACAAGCGACCCCATCAAAAACAAAAGCCAAAATTGACAAATGGGATCTAATTAAACTAAAGAGCTTCTG
CACAGCAAAAGAACTACCATCAGAGTCAACAGGCAACCTACAAAATGGGAGAAAATTTTGTCAACCTACTCATCTGAC
AAAGGGCTAATATCCAGTATCTACAATGAACTCAAACAAATTTACAAGAAAAAAACAAGCGACCCCATCAAAAAGTGGG
TGAAGGACATGAACAGACACTTCTCAAAGAAGACATTTATGCAGCCAAAAAACACATGAAAAAATGCTCACCATCACT
GGCCATCAGAGAAATGCAAATCAAACCACAATGAGATACCATCTCACACCAGTTAGAATGGCGATCATTA AAAAGTCA
GGAAACAACAGGTGCTGGGGAGGATGTGGAGAAATAGGAAAAGTTTATACTGTTGGTGGCACTGTAACTAGTTCAAC
CATTGTGGAAGTCAGTGTGGCGATTCTTCAGGGATCTAGAATAAGAAATACCATTTGACCCAGCCATCCCATTACTGGG
TATATACCCAAAGGATTATAAATCATGCTGCTATAAAGACACATGCACATGTATGTTTATTGCAGCACTATTCACAATA
GCAAAGACTTGAACCAACCCCAAATGTCCAACAATGATAGACTGGATTAAAGAAAATGTGGCACATATATACCATGGAAT
ACTATGCAGCCACAAAAAATGATGAGTTCATGTCCTTTGTAGGGACATGGATGAAATTGGAAATCATCATTCTCAGTAA
ACTATCACAAGAACGAAAAACCAAACACCGCATATTTCTAACTCATAGGTGGGAATTGAACAATGAGAACACATGAACAC
AGAAAGGGGAACATCACACTCTGGGGACTGTTGTGGGGTGGAGGGATGGGGGAGGGATAGCTTTAGGAGATATACCTAA
TGCTAAATGATGAGTTAATAGGTGCAGCACACCAGCATGGCACATGTATACATATGTAACCTAACCTGCACATTATGCAC
ATGTACCCTAAAACCTTAAAGTATAATAATAAAGTAAAAATAAAAAAAGAAAGTTTACATATAAAAAATAAATACATA
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TTCAAGAGATGGCCTGACTGCCTCTAAAAGCCTAATTCATTTGCATAAAACAAAGAAATGACCTGAACTAGAACTTGTA
TTTAAAGGGGAAGCAGAGCATAAAAGTTTGGAAAATTTTCAGCCAGACCATGCAGTAGAAAAGAAAAACAATTTTCTAG
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GACCAGGCCCAAGACCCCGCTGCTCTCTGCAGCCTTGGGACATGGTGCCCTGCATCGCAGCTGCTCCAGTTCCAGCTGT
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TTGGGCCTGTGAGTGTGCAAAAGGTAAGCGTTGAGGTTTGGGAACCTTTGCCTAGATTTTCAGAGGATGTATGGAACTC
CTGGATGTCCAGGCAGAAGTGTGCTGCAGGGGTGGAGCCCTCATGGAGACCCTCTTCTAGGGCAGTGCAGAGGGGAAAT
GGGAAATGTGGGGTTGGAGCCCCCACACAGAGTCCCTCCTGGGGCACTGCCATAATGGAGCTGTGAGAACGGGGCCACCA
TACTTCAGACCTCAGAATGGCAGATCCACTGACAGCTTGCTCTGTGTGCCTGGAAAAGCTGTAGGCACTCAACACCAGC
CTGTGAAAGCAGCCATGGGGGCTGTATTATGCAGAGCCACAGGGGTGGAGCTGCCCAAGGACTTGAGCCCCTGCTTGC
ATCAGTATAACCTGGATGTGAGACATGAAGTCAAAGGAGATAATTTTGAAGCTTTGAGATTTAATGACTCCCCTGCTGG
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TCTCTTCAAAGAGTCTCATCTTTGTCTCTCATAAAAGATGAGACTTTGGACCTACACTTTTGAGTTAATGCTAGAAATGA
GTTAAGACTGTTGGGAAGGCATAATTGTGTTTTGAAATATAAGAAGGATATGTGATTTGGGAGGGGCCAGGACAGAATT
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GGTAATTAGATCATGGGGGCTGATTTCCCCCTTGCCATTCTCATGAGACCTGGTTGTTTAAAGTATGTAGCACTTCCC
CCTTTGCTCACTCTCCTGCTCCACTATGGTAAGACATGCTTGCTTCCCCCTTACCTCTGCCATGATTATAAGTTTCCCTG
AGGCCTCCCAGCCATGCTTCCTGTACAGCCTGCAGGACTGTGAGTCAATTAAACTTTCTTCTTATACCATTTGATCCA
GCAATCTCATTACTGGGTATATACCCAAAGGATTATAAATCATTCTACTATAAAGACACATGCATACGTATGTTTATTG
CAACACTATTTACAATAGCAGAGACATGGAACCAACCTAAATGCCCATCAATGATAGACTGGATAAAGAAAATGTGGTA

Fig. 9.13

Fig. 9.14

CCTGGAATTGCTCGTTAAGACACACATTGCTGGGATCCACCTCCAGGGTGTGAGAGTCAAGTCTGGGGTGAAGCCTGAG
AATGTGCATGTCTAACCAGTTCCAGGTGATGCTGATGTTGCTGGTTTTGAGACTATATTTTGAGATTAAGAACTGCCT
TGGATTAAGAGCTGACCTGCGATCTACAGCTCAAATAGTGAAGTAAACATCCTAAAGAAAATGGAAAAACCAGTGCAGT
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TATATATAACTCTTGCCACCTAGGATGTTTCATAGTAACATAAGACAGTATTTTGTGTTTCCAAGTAATTTTAATGATC
CTGTAGATCCTCTTCCTTTTTGTATATAATATCAATCTAATAGTTTCTTGTTTAAATATAAAAATGAAATCTTATTTTAC
ACAAGCAGCAAGCAGCTATTTTTTTCAGATTTTCCCCCTATAATCTAAGGGAAAGTTATTTTAAAATAGAAAAGATGTGG
GCTTCAAAAAAAGCTTTGCAAATATGTTGCAATAATACGAATGATTTTCAGTGTGAAATCCATTTGTGAAAGCAGGCTT
TGCTTATATTTTGGGTCTGCTTCTATAAAAATGCTCAGATTTGCTTTTATTAAGATCATACACTCAGTGACCTGAGGA
CCAGATGGAGGTTATAAGCAGCTCTTTAAGGCTTCAGAGCTTAGCCTAGAGAGTCAAACAGCTCTTTGAACTGGCGTCT
CAGCTCTGTCAATTAGGCAAATGTTTTATTTTCTGCAAAGAACCAGCCTTAGGCTTTGTTGATCTACGTATGTATATTT
GTTTTTTTATTTATTTGCATTATTTCAATTCATTCTGTTTTCTTTGACTTTAATTTGCTAGTTTTCTTTGAAGCTTGTGAG
ATAAATGCTTAGATCATTGATTATCAAAATTGTGTCTATTCTAATATATATATAATATATATATGATAAGGCTATACAT
TTTCCCCTTAGCACAGCTTTAGTAGCCTCCAACAAATTTTGATGTGCTGTGTTTTTATTATAAATCAGTTTGAAATATA
TATTCTAATTTTATTATGATTTCTTGAGCCTATGGATTATATAGAAATATATTTCTTAAATTGTAAACATATGGGGATT
ATTTGATGTCTCTAAGCCGCAGATTTCCCTATTTTATTTTATTTATTTTATTTTATTATTATACTTTAAGTTTTAGGGT
ACATGTGCACAATGTGCAGGTTTGTACATATGTATACATGTGCCATGATGGTGTGCTGCACCCATTAACCTCGTCATTT
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TCCTGTGTCCATGTGTTCTCATTTGTTTCAGTTCCCACCTATGAGTGAGAACATGCGGTGTTTTGGTTTTTTGTCTTGCAA
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GGCTTGGTTTGAAGTCTTTGCTATTGTGAATAGTGCTGCAATAAACATATGTGTGCATGTGTCTTTATAGCAGCATGAT
TTATAATCCTTTGGGTATATACCCAGTAATGGGATGGCTGGGTCAAATGGCATTTCTAGTTCTAGATCCTTAAGGAATC
GCCACACTGATTTCCACAATGGTTGAAGTAGTTTACAGTCCCACCAACAGTGTGAAAGTGTTCCTATTTCTCCACATCC
TCTCCAGCACCTGTTGTTTCCCTGACTTTTTTAATGATTGCCATTCTAACTGGTGTGAGATGGTATCTCATTTGTGGTTTTG
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GATTGTGGATATTAGCCCTTTGTCAGATGAGTGGGTGCAAAAATTTTCTCCCATTCAGTAGGTTGCCTGTTGACTCTG
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GGTGTTTTAGACATGACGTCCTTGCCCATGCCATATGTCTGAATGGTATTGCCTAGGTTTTCTTCTAGGGCTTTTATGG
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GGTTTTGTCAAAGATCAGAAAGTTGTAGATATGCAGCATTATTTCTGAGGGCTCTGTTCTGTTCCATTGGTCTATATCTC
TGTTTTGGTACCAATACCATGCTATTTTGGTTACTGTAGCCTTGTAGTATAGTTTGAAGTCAGGTAGCATGATGCCTCC
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TCCATTTCTGTGAGGAAAGTCATTGGTAGCTCGATGGGGATGGCATTGAATCTATAAATTACCTTCGGCAGTGTGGCCG
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CAGTGGTTTTGTAGTTCTCTTGAAGAGGTCCTTCACATCCCTTGTGAGTTGAATTCCTAGGTATTTTATTCTCTTTGAA
GCAATTGTGAATGGGAGTTCACTGATGATTTGGCTCTCCGTTTGTATTGGTGTATAAGAATGCCTGTGATTTTTGAC
ATTGATTTTGTATCCTGAGACTTTTCTGAAGTTGTTTATGAGCTTAAGGAGATTTTGGTCTGAGACGATGGGGTTTTCT
AGATATACAATCATGTCTGCTAACAGGGACAATTTGACTTCTCTTTTCTAATTTGAATGCCCTTTATTTCTTCT
CCTGCCTGATTGCCATGGCTAGAACTTCCAACACTATGTTGAATAGGAGTGGTGAGAGAGGGCATCCCTCTGTCTTG
CCAGTTTTCAAAGGGAATGCTTCCAGTTTTTGTCCATTCCGGTATGATATTGGCTGTGGGTTTTGTCATACATAGCTCTTA
TTATTTTGAGATACGTCCCATCAATACCTAATTTATTGAGAGTTTTTAGCATGAAGGGTTGTTGAATTCTGTCAAAGGC
CTTTTCTGCATCTATTGAGATAATCATGTGGTTTTTGTCTTTGATTCTGTTTATATGCTGGATTACGTTTATTGATTTT
CATATGTTGAACCAGCCTTGCAATCCAGGGATGAAGCCCACTTGATCATGGTGGATAAGCTTTTTGATGTGCTGCTTGA
TTCGGTTTTGCCAGCATTTTATTGAGGATTTTTGCATCAGTGTTTCATCAAGGATATTGGTGTAAAATTCTCTTTTTTGT
TGTGTCTCTGCCAGGCTTTGGTATCAGGATGATGCTGGCCTCATAAAAATGAGTTAGGGAGGATGCCCTCTTTTTCTATT
GATTGGAATATTTTCAGAAGGAATGGTACCAGCTCCTCCTTGTACCTCTGGTAGAATTCGGCTGTGAATGCGTCTGGTC
CTGGACTTTTTTGGTTGGTAAGCTATTATTATTGCCTCAATATCAGAGTCTGTTTTTGGTCTTTTCAGAGATTCAACT
TCTTCTGATTTAGTCTTGGGAGGGTGTATGTGTCCAGGAATTTATCCATTTTTTTCTAGATTTTCTAGTTTATTTGTG
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TTGCATCTATTTGATTCTTCTCTCTTTTTCTTCTTTATTAGTCTTGCTAGCGGTCTATCAATTTGTTGATCTTTTCAAAA
AACCAGCCTCTGGATTCATTGATTTTTTGAAGGGTTTTTTGTGTGCTATTTCCCTTCAGTTCTGCTCTGATCTTAGTTA
TTTCTTGCCCTCTGCTAGCTTTTGAATGTGTTTGTCTTGTCTTCTAGTTCTTTTAAATTGTGATGTTAGGGTGTCAAT
TTTAGATCTTTCTGCTTTCTCTTGTGGGCATTTAGTGCTATAAATTTCCCTCTACACACTGCTTTGAATGTGTACAG
AGATTCTGGTATGTTGTGCTTTGTTCTTATTGGTTTCAAAGAACATCTTTATTTCTGCCTTCATTTTGTATTGTACTC
AGTAGTCATTCAGGAGCAGGTTGTTTCAGTTTCCATGTAGTTGAGCGGTTTTGGGTGAGTTTCTTAATCCTGAGTTCTAG
TTTGATTGCACTGCGGTCTGAGAGACAGTTTGTCAATAATTTCTGTTCTTTTACATTTGCTGAGGAGAGCTTTACTTCCA
ACTATGTGGTCAATTTTGAATAGGTGTGATGTGGTGCTGAGAAGAATGTATATTTCTTTTGATTTGGGGTGGAGAGTTC
TG TAGATGTCTATTAGGTCTGCTTGGTGCAAGAGCTGAGTTCACTTCCCTGGGTATCCTTGTTAACTTTCTGTCTCATGGA

Fig. 9.15

TCTGTCTAATGTTGACAGTGGGGTGTTAAAGTCTCCCATTTATTATTGTGTGGGAGTCTTAGTCTGTTTGTAGGTCTCTA
AGGACTTGTTTTATGAATCTGGGTGCTCCTGTATTGGGTGCATATATATTTAGGATAGTTAGCTTTTCTTGTGAATTG
ATCCCTTTACCATTATGTAATGGCCTTCTTTGTCTCTTTTGGTCTTTGTTGGTTTAAAGTCTGTTTTATCAGAGACTAG
GGTTGCAACCTGTGACTGTTTTTGTGTTTTCCATGTGCTTGGTAGATCTTCCCTCCATCCCTTATTTTGAGCCTATGTGTGT
CTCTGCACATGAGATGGGTCTCCTGAATACAGCACACTGATGGGTCTTGACTCTTTTTCCAATTTGCCAGTCTGTGTCT
TTAATTGGAGCACTTAGCCCATTTACATTTAAGGTTAATATTGTTATGTGTGAATTTGATCCTATCATTATGATGTCA
CCTGGTTATTTTGCTCGTTAGTTGATGCAGTTTCTTCCCTAGCCTTGATGGTCTTTACAATTTGGCATGTTTTTGCAAGTG
GCTGGTACCGGTTGTTCCCTTTCCATGTTTAGTGCTTCCCTTCAGGAGCTCTTTTAGGGCAGGACTGGTGGTGACAAAATC
TCTCAGCATTGCTTATATGTAAAGTATTTTATTTCTCCTTCACTTATGAAGCTTAGTTTGGCTGGATATGAAATTCCTG
GGTTGAAAATCCTTTTCTTTAAGAATGTTCAATATTGGCCCCACTCTCTTCTGGGTTGTAGAGTTTCTGCAGAGATAT
CCGCTATTAGTCTGATGGGCTTCCCTTTGTGGGTAAACCCGATGTTTGTCTCTGGCTACCCCTTAACATTTTCTCCTTCAT
TTCAACTTTGGTGAATCTGACAACTATGTGTCTTGGAGTTGCTCTTCTCGAGGAGTATCTTTGTGGCATTCTCTGTATT
TCCTGAATTTGAATGTTGGCCTGCCTTGCTAGATTGGGGAAGTTCTCCTGGATAATATTCTGCAGAGTGCTTTCCAACCT
TGGTTCCATTCTCCCCGTCACTTTCAGGTACACCAATGAGACGTAGATTTGGTCTTTTCACATAGTCTCATATTTCTTG
GAGGCTTTGTTCAATTTCTTTTACTCTTTTTTCTCTAACTTCTCAATTCATTTCAATTCATTTCACTTCCATTACTGA
TACCCTTTCTTCCAGTTGATCGAATCGGCTACTGAAGCTTGTGGATGCATCACTTAGTTCTCGTGCCATGGTTTTTCAGC
TCCATCAGTTCAATTAAGGACTTTTCTACACTGGTTATTCTAGTTAGCCATTCGTCTAATCTGTTTTCAAGGTTTTTAG
CTTCTTTGCGATGTGTTTGAACCTTCTTCTTTAGCTTGGAGAAGTTTGATCATGTGAAGCCTTCTTCTCTCAACTTGTC
AAAGTCAATCTCCATCCAGCTTTGTTCTATTGCTGGTGAGGAGCTGCATTCCTTTGCAGGGGGAGAGGTGCTGTGATTT
TTAGAATTTTCAGCTTTTCTGCTCTGTTTTTTCCCATCTTTGTGGTTTTATTTACCTTTGGTCTTTGATGATGGTTAC
GTACAGATGGGGGTTTTGTTGTGGATGTCCTTTCTGTTTGTAGTTTTCTTTTAAACAGTGAGGACCCCTCAGCGGCAGG
TCTGTTGGAGTTTGCTGGAGGTCCACTCTGGACCCTGTTTGCCTGATATTACCAGCAGAGGCTGCAGAACAGCGAATAT
TGCTGAACAGCGAATATTCTGAACAGCAAATATTGCTGTCTGGTAGTTCTCTGGAAGCCTCATCTCAGGTGGGTACC
TGGCCATGTGAGGTGTCAGTCTGTCCCTACTTGGGGGTGCCTCCCAGTTAGGCTACTTGGGTGTCAGGGACCCACTTGA
GGATGCAGTCTGTCCGTTCTCAGATCTCAGACTCCTTGCTGGGAGAACCCTACTCTTCAAAGCTGTCAGACAGGGA
CATTTAAGTCTGCAGAGGTTTCTGCTGCCTTTTGTGTTGGCTATGCCCTGCCCCCAGAGTTGGAGTCTATAGACGCAGGC
AGGCCTCCTTGAGCTGAGATGGGCTCCACCCAGTTTCGAGCTTCCCAGCCACTTTGTTTACCTACTCAAGCCTCAGCAAT
GGTGGGGCGCCCCCTCCCCAACCCTTGCTGCTGCCTTGCAAGTTCGATCTCAGACTGCTGTGCTAGCAATGAGCGAGGCTC
CGTGGGCGTGGGACTCTCCGAGCCAGGCNCGGGATATAATCTCCTGATATGCCGTTTGCTAAGACCATTTGAAAAGTG
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CCTGCAGTGTAGCAAAGTTTGAAATGTAGATAGCCAGAAGTCAGTCTGGGGGAAAGCAATTTCCAATTTTCAAGCTTGTA
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ACAAATTCACTATACCATTTTTTTTCTTTTTTGTGCTGGAACATGTATAAAATAGAATATATCTGAATTTCCATGTAGGGCT
GAAACATGCTGTCAATAGAAACAATATGCCTACATCTCCGTCTCTCTTTCTGTGTGTGTGTGTATGTATGTGTGAAG
TCATATGTTTTATGCTTTTCTATATATCAGATTATGTTTTAGCATTTTCAGAGGCACTGTGTCTGCTAAAATCCTGTGT
TTCCAGATGAAATGGCAAACATTATTTCCAGCAATGTGATAAAACAACCAAAGAAAGTGTTCAGTGACCCGATTCATA
AAATTCACCAATAAGTCAATATATGCAGTAAAATAGTTATTACAAGAGTAGCTAAGAGTAATTAATATGAGTAAGTCTG
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ACCAAATCCAGCCTACTGTTTTATTTGTAAAAAATTACATTTGAACACAGTCTTACCCATTTATTAACCTTACTGTTTTAT
CACTGCTTTTGTCTAGAAAAGCAGAGTTGAGTAGTTGAAAAAAAGACTGTGGCCAGGCACAATGTCTCATACCATAAT
CCCAGCACTTTTGGGAGGCCAAGGTGGATCACTAACTGGTCACTAACTCTTGGGCTCAAGTGGGTGGATTACTTTAGCCC
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GAACTGAGATCATGCCACTGCACCTCAGCCTGAGTGACAAAGTGAGATCCTATCTCCAAAAAAGAAAAAAGAAAGA
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TGGAAGAAAGAGGCTAGCAGAAGCTTCAAAAATATCAAAAACCTCTCTTACTGTGTGGCAATATAAACTAAAACTGAT
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AGATGTTTATGAGAACCAGGGTCTTTTCAAACAGGGGGTGAAAAAGGAGTGAGAAGATGTGCATGGCAGTGAAGAAGAA
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AAGTACTGTCATCTTTTGTATGAGCCATATTTCAATTCACAGTGTTTAAATGACTTTTTTCCAATCATTACTCCATAA
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TAACACAATATTTTATTTTCTAAAGATCTTGCAAGTGTGTCATCTTTATCACCTTTTAAAGTTTGCATTTTATTTGT
CACTTTTAAATTTTAAATAAATGTCCATTTTGTATCTATCATTTCTGTGTCTTTATGTAATAGTATCTAATATTCTAT
ATCTGTATCAAATATTCTGTATGTACTTCATATTCATTTTATCTTATTTTATTTATTTATTTATTTATTTATTTA
TTTATTTTAGACAGAGTCTTACTCTGTCTTCCAGGCTGGAGTGCAGTAGCACAATCTTGGCTCACCACAACCTCCACCT
CCCGGGTTCAAGCAATTACCCTGCCTCAGCCTCCTGAGTAGCTGGGGTTACAGGCACACGCTACCATGTCCAGGTAATT
TTTGTATTTTATAGTAGAGATGGGTTTTGCCATGTTGACCAGGCTGGTCTCAAACCTCCTGACCTCAAGTGATCTGCCCC

Fig. 9.16

CTTGGCTTCCCAAAGTGCTGGAATTACAGATATGAGCCACCACATCCAGCCCCATTTTACCTTAATTTGTAAGAATATA
TAATTATATTTATAATTTTATTCATTTTATATTAATTTTCTATTTCTTAAAGGCTAAATCTCTAGTTTCCTTTTTTATT
TTTCAGGTAGAGGATTTTAATTCACCTGAGTACCTGATCATTTTACATACCTAGTCAAATAAATATGAAATTTGAAGAAC
TAAGATTAAGTTTTTAAAAATTCTTAAGTGCATACAATCCATTGAAGAAAACAAAAATTGCTTAAGCAGTTTTTAAAACC
TTGATAAGAGGTCATCAACATGATTTAAATGTGAAATTAGCTAATGTAAGTAGCAGTTAGAATGAGATTTAGCTTATAT
ATTTGGAGGTATAATGGAAAACCTCTGGTGAGGACTCTTTTATTGACACCAAGAGAAACATGAGAAAGAAAAATGAGTT
GCATTTTTCTGCATTTTTCCAGAAGTGAAATGCTTTGCAACATCCTCAGAAAGAAAGAGAATACATGAGTAATATGAAA
GAAAACATATATAAAGGATTTAATTATTTCCAAATTGCCTGGGTGTTTTTTTTTAGTGCTGCTTTTTTTTTTCAAGCAGGAA
TAAGTGTAGAACTTTTATTTATTTATTTTTTAAAGGAATGGTTACTATTCTTAGGAAAAGTGGCAGTAAATATAGTTAA
TAACGGTAATTATAATTTTATAAACTCATTCAAAGTCTTGATTTTAAAGGCGATAGTAAAAAAAATATATATCTATTT
ATCTTGTTCCCTGAAAAATGGTAGCACTCCTGACCCTGAGAGACTGTTCTCCTCGGCATAATTGATGGCCTTCAAAGC
CATCATACTTCAGTATTCATGTCAAAGATTTATTTCCATTTGCAAAATTTGCTTAGAACTCACCTATATTTACCTTTC
CTCTTATGACTTATAAGATAATATTAACATCTACATAACAGAATCTCACTCACAGAATGTTTGAGCCACGAAGGTGCTC
TGAGACCCTCTGCCTCAACAGCCTCATTATAGATGAGAGAAGCCAGAGAGGTGAGGACGCCTTCCAAAGCTCTAGCA
CTTGTTTTCTAGACATGGAACCAGGACAGCCCTTGGCATTTCCCACCACATTTTCTGCTTATTATGTGGCATGAGCTTG
TCTTTAACACTGAATTAGAATTATTGCACATTACATTGCATTCATCAAGGCCTACCCTTCAGGCAGTCTGATGTAACAG
TCTATTATAAGACATCCGAGAAAAACCATCCACTATCCTTTCAGATGTTCTCAAACAATTTGTATTAAAGAGCTAATAA
AATGAAACAGAAAATGCCAGGTAACTTTTTTAAATAAAGATTTTGTGATTCATTGGGTTTAATTATTGGTTTTATTTT
AGCATTACTTTGATACATTTTCATGAAGTATACAGACAGAACTAAAGCCACATTTAAAAGGCAATTAATAAAACCCAGAC
ATTATGATATAATTTACATACATTTAGATATATTTAAAGTGTACAGTTTCAATTTATTAGTATAACCAAGAGTTGTGCA
AACATCACCACCATCTAATTCAGAACATGTTAGAACCCCAAAGAACTTCATATTCATTAGCATGTTGCATTCCTGG
CAGCTCCCTCTTGCCACCCTGTACCCTCAGTCTTAGGCAACCCTAAACAACCTTTCTGCTCCTCATAGACTTGTCTATTC
TAGACATTTCTTATAAATGGCATCATACGATATGTGGTATTTTGTGACTGGCTCTGGCTTCTTTCACTTAGTATAATGC
TTTCACATTTATCATGTAGCATTGTGTAAGTGTCTTTCCCTTTTATAGCTGAGTAATATTCATTGTATGTGTATTCCAC
ATCTTTATTTATCCATTTCATCACTTGATGAATATTTGGATGGTTTTCACTTTTTGGCTACTATGAACAATGCTGCCATG
AACATTTGTGAACAAGTTTTTAGTGTGGCATTATGTTTCCATTTCTTTTGGGTACATATCTAGAAATAGAATTACTGGGT
CATAGGGTGACTCTATGTTTAATATTTTGAGGAATTCCAGACTGTTTGCAAAGTGACTGCATCATCTTACAATCTCAC
CAGTAAGTATGGCATGAGGATTCCAATTTCTTCACAGCCTTGTAATAAGGAAATTTTACAATTATAATTATTTGGCAA
AAATTCAATGAATTTTTTGAGCATCTATTTTGTGAAAGGTCTTTTTTTGAGGTGCTATAATTTTTTAACACTATCAATTT
TTGTGGAAAAAATTAGAATTTTCTAAAGGACATTATTTTAAAGTTTATTACTTATTATTTATTAATTTTTTATTAATA
CATAATATTGCTATGTATCAGGTACATGTGATAGTTTGATACATGCATACAGTATATAATGATCAAATCAGCATATTTA
GAAAATCCATTACCTCAAGCATTATTTCTTTGTGTTTGGAACATTTTCAAGACTTCTCTTTTCAAGCTATTTTGAAATATG
CAATATATTTTTTTGTAACTATAGACACTCTATTGTGCTATTAAACACTAGAACTTATTTCTTCCACATAACTGTATGT
TTGTACCCATTAACCAATCTGTCCTCATCCCCCAGCCCTTTCCAGCCTCTGGTATCTATCATTCTATTCCCTACCTC
CATGAGGTCAACTTTTTTAGCTCCCACATATGAGTGGGAACATGTGATACTTGTTTTCTGTGCCTGGTTTATTTCACTA
AACATAATGACTTGCAATTCATCCCTGTTGCCGCATATGATGAGATTTCAATTTTTAATGGCTGAATAGTATTTTGTG
TGTATATATAACCATATTTTCTTTATTCATTTCATCTGTTGATGGACACTTAGGTTGATTTTACATACCTTGGCTGTTGTGAA
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GGGATTGCTGGGTCAGATGGTAGTTCTATTTTTTAGTTTTTAAAGACACTTTCATACTGTTTTCCATAGTCGTTGTACTA
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CATAGAAATCTTTCACCTCCTTGGCTAAATTTATCCCTGAGACTTTTTTTGAAGTTATAAATGAGGTTGCTTTCTTGGTT
TCTTTTTTCAAGATAGTTGGTTATTGGTGTATAAAAAACACAACCGATTTTTTATATTGATTTTGTGTCTTGTAACTACT
GAATTTGTAAATCATGTTCATCTGAAGTAGAGACAATTTGGCTTCTTCTTTTTTTTTTTTTTTGAATTGGAATCTCTCT
CTGTTCATCCAGGCTGGAGTGCTCTGGTGTGATCTTGACTCACTGCATCCTCTATTTCCCAGGCTCAAGTGATCCTCCCA
CCTCAGCCTCCTGAGTAGTTGGGACTACAGGTGTGTGCCACCACACCTGGCTAATTTTTTGTATTTTTTCATAGAGACAGG
GTTTCATCATGTTGTCTAGTCTGGATTTGAACTCCTGGGTTCAAGCAATCTGCCCACCTCAGCCTCCCGAAGTGCTGGG
ATTACAGGCTTGAGCCACCACAACCTGGCTGGCATCCTGTTTTCCAGTTTGATGCTTTTCAATTTCTTTCTTTCCGGA
AAGAGAAAGTCTGGCTAGGACTTCCAGTATAATGCCGAATAAGAGTGCTTAGAGTAGGTGTCCTTGTCTTATTCTAGTT

Fig. 9.17

CTTAGAGGAAAGGCTTTCAGTTATTCCCCATTCAGTATGATGTTAGCTGTGGGCTCGTCATATATGGCCTTTATTATGT
CAAGGTATGTTCTGCTATACCTAATTTGTTGAGAGTGTTTCATCATGAAGGGCAGGGTGAAAGGGATTCTTTTCTGAAG
GAGGTTGTGTCAGTGTCTGATGAGGATATAGAAGTCAGTGTTTGGTGAGAGATAGTAATTTGAAGAGGGGAACATT
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CAAACATAGGGAAGAGCTGCTACTTCTAGGACTGAGGGAGGATACCCAAGAAAAGAACAAACATGGAAGGGTCTTTCAC
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CAGGCCAGGCCTGGTGAGTAGGGTACTGGCTGTTGGGTGCCAGTGGATCAGCACTGTGGTCAAAAAAGTGCCTTCTAGG
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ACAATAAACTCAGTAGGAAGCCCTCACTAGGTGCTGGTGGAACCTCACTGTAGGGTGACTCTCCCATACACCACTGGTG
GCGGCCACAGGTAACAAGAACCAGGAAGAATCAAGAAGGAATGCTCCTTTATTTGCTATACCTTGTTTTATTTTTATTTT
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CCTCCACCTCCCAGGTTCAAGTGATTCTCCTGCCCTCAGCCTCCCGAGTAGCTGGGATTACAGGTGCGTGCTAATTTTGT
ATTTTCAGTAGAGATGGGGTTTTACCATGTTGTCCAGGCTGGTCTCAAACCTCCTGGCCTCAGGTGATCCTCCTGCCTCT
GCCTCCCAAAGGGCTGTGATTACAGGCATGAGACACCGCATCTGACCCCTTTACTATACCTTGAGTGCTCTCCCTTACA
CTCTACCAGCAACAGATGACATTGCACTGGCTGACCGAGGAGCCAGATTAGTATCGTGGAACAGGGGCAAAGAAGGGTGG
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AATGAGGAGACACACAGTTCCAACAGTTATCGTAGTCTCATCTGGCTGTCTTAAATACTCCTCAAATCAAAGTCCCACT
GAATATTCTCTTACCTAAAGGCTAAATTGTAGAGTTTATATTCAACAACTTTTATAAAATAATGAAGAGAGAAAAAGGA
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CAAGAATTTATACACAAAGGTGAAAGGCTTTTTGAAACAATCAGAGTTTTCTTTCAAGTCTAACTTCCTGAGTAATTTA
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CCCATCTAAAAAAGATGATAGTTTACATTGTTTTGAGAATTGTGGAGTGGGAGAAGCAGACCAACCTTTCATACTGCT
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GCTTAATCTGTGGTGGCATGAAATGATGAAAAAATAAATATATTTTCTCTTTTTTTTTTCTGTTTTTTATTCTCCATCCA
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AGGATGTTGGAGTGCTAAGATCCCTGAATGTGGTACCTGTAGATAAATACAGTGACTTTGAGACGCAGAGCAAAGACTC
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GCCAGTATGATTTGAGCCAGAAGATTAAGTTCCATAAAAAGTGAAGAAAAGTCAACAATTTAACACAAGAAATGTTTAC
CGACTAACTATTCTATGCCAGGTCTGGGAGTATGAGGAAATAAATGTGACATATCCCTTGAGAAAGCAGAAATAATCCA
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TATACCCCTGAATATCATTACTTGAAGAAAAAATACACATTTCTTTTGAAGAAAGTAAAATCAGAAATCTTGAGGGTT
CTCTCGAAGAAAATCAGACCTGTAGTTCTAACCAGTCAACTGACCTTAGTTGAGAATGTTAATGTGCCAGAGAAACAAA
TGAATGCTCACCTTTCCGAGGAAGACAGTAGAAAATAACCCGCTTGACTATCTTCAGCTGGAAAAAAGAACTCTTTTTT
TTTTTTTTTCTGTGACGGAGTTTCACTCTTGTGTTGCCAGCCTGGAGTGCAATGGCACAATCTCAGCTCACTGCAACCTC
TGCTCCCGGGTTT CAGGCAATTCTCCTGCCCTCAGCCTCCCGAGTGGCTGGGATTACAGGCGTGCAACCACCGCCCGGC
TAATTTTTTGCATTTTTTCAGTAGAGACAGGCTTTCACCATGTTAGTCAGGCTGGTCTCGAACTCCTGACCTCAGATGATC

Fig. 9.18

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CACCCACCGCAGCCTCCCAAATGCTGGGATTACAGGCGTGAGCCACCGTGCCCATTCGAAAAAAGAACTCTTTAATCC
ACTATACCTAACCTTTTACTCTCTCCTTTAGATTTAAATGATTAACCTTCTGATAATGCTGCAGACATTTTTTTTCTGA
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CATGTTCTGTTTCAATGTTGTGAACATTCATTTTAAACCTACTATATTTAGCAGAGGTTTAATTACCCAAATTGGGAAA
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AATAAGAGTGAAAAATTAAGTTGGCATGATAACAAAATAGACTATGTAGTGGAATATCAAAAGTAAGAGAAAGGAATCC
TAACATAAATGAGAGTACAACATTTCAACATATTACCATAAAAAGTTTAAATTTGTCTAATTCACCTATTGACTCAAATT

Fig. 9.19

[illegible]

Fig. 9.20

AGTGAAAGGGTGAAGGTGAAGTGAACACTAGAAACAGAGAGCAAATGGAGGCTGAACCTTATCCTTTTATCAGGAACC
CACTCATGGGGCTCTGCCCTCGTGATCTAATCACCTCTTAAAGGGCCACCTCCCAATACCATTACATTAGCAATTAAA
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CTCCAAGAGAAAAGGAATATTCTCTTTTCTTCTGAAAAAGCACTACAGTTTATTGGTCAGCACCAAGTGTGTTAACTT
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GCAACAGAAGTGGCTAAACAGAATAATTCCAGGCTTGAAGTTCCTTTTTTCATCTGGTTCCCTCTGGGAAAAGGCTATGA
AAGCTAAGGGTGGTGGCCAAGTCCCAAAGGCCAAGGTCACCATCAAACATTTTCTGAGAATCTACCAAAAACACAGGTC
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AATCTTTTTTGCTTAAGTCCCTTTGTGCTGCCACATCAATTTGATTTCCACAGCAAAATTATTATAAGCCTGCATTATC
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ACCGTGTTAGCCAGGATGGTCTCAATCTCCTGACCTCGTGATCTGCCCCGCTCAGCCTCCCAAAGTGCTGGGATTACAG
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GGTGGGAGGAGGGGAGGCTCAGGCATGGCAGGCTGCAGGTCCCGAGCCCTGCCCTGCGGGGAGGCAGCTAAGGCCTGGT
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CAGTGACAGTGGCGGGCTGAAGGGCTCCTCAAGCACGGCCAGAGTGGGCGCCAAGGCCGAGGAGGCACCGAGAGTGAGCG
AGGGCTGCAAGGGCTGCCAGCACGCTGTACCTCTCAATACGATCTGTGAACATGTGCCCTTTCTCAGTTTCAAGAGCC
ATTTAGAGCAAATGTACTTATTTACTTTTATCCAGGCTCTGCTTACTCCTCTCCTCTCCACAGCCATCTCCTCCATTAGA
AATGCTTTTCCCTGCCATTTTTTGACAGACCCTGAGCCTCCTGGCCTCCCTGCCCTGAGACGCATTATAGTCTCTCTGACC
TGGACCACTGCATGCAGCTTGGGAGCCTCCCTGTCTTTAAGGTATCGGACAAAATATAGATGAGAGTAGCAAGGCATTA
GGAGTAGTAGTAATAGCAGTACAAACAAGAATAATAATTTTTATGATAATAGTAGCTAACATTTGTTGAACTCCTTTTA
TGTGCCAAGCATTTTTATGTTTATTTATCTTACCACCCTTGACAAGGACCTTATGAGGTGGGTACTATCAACCCAGTTTCT
CAGAAGGGGAACTGAGGCTTATAAAGAATATACAACTTGCTCCAAGTGGTAACTAGTGGAGAGAGGATTTAAACCTTG
GCTCTTCAACATCCATAACCTATGCATTTAGTTGCTGTTCATAAATAGTCACCTGCATATGTTTATCTTTTACC
TGTGTTTCAATCCATCTTTTTTTGTGAACTCCTGGAAGGCAGGTGCTGTCTTCTTTTCAATCTTTTATACCATCTCAGAGG
GCTGTAACAATACATGCTCAGAAAACATTTCTGTTCAACCTGCATGAAGTACAAGGCAGGCTGAAATGTAAAGAAATAC
ATAAATTACACTGAGGAAGAGCAAATCCTTTTCAATATAACTTTTTCTTGGCATTACCTCTTAGGGGAAAAGAATTTCTCAA
GCAAGCTGTAAGTGAAGTGGTTACCAAAATCAGAATGAATTTCTCCCAATCACGGGGGACATTCCATAAGCCTTTGGAG
TGAGGTCTATTTTAAGGATTTTGTTCAGTTCTCTTGTTCCTGGCCACACTGCCAGCACTTCAATTGGCTAA

Fig. 9.21

TAGGGATTCTGAATAAGAAAGATACAGGCAGCTCACTTGGAAAGTTACGTTAAGGACTAATAAAGATCTTGGAGTGATTT
TCAACAAGATAAGATCAGCATCCTGGTATTGAGTTATTTTGCTACATTAATTAATTTTCAGTTTGTCTAATGGTCTTTG
GCCTCCTGCTTAAATGTCAATTGAGGCAACAGATTAATTTCCCTTAAATAGTGGCCTGTCATGTCTTGTACTTAGTTT
TGGGTTTTATAAAGTAAGAACTAAGATGGATATTTATCAATCGAGCAGATTTCTGTGTTGTGTCCTCTAAAGCCCTTGG
GGTCAGTGTCTCACTGTAGCATCACTAACACCTTTTATGGAATCAAGTGATTAATAAATGTGTCCTCATGAGACTGTGA
TGCTTCTGGGTGTTCAATTTTAGATACCTTAATAATTAGTCTCAAGAGTAGCTCAGTTATTAAGTGATTTTGGGGTTGCA
TCATGTCAGGGAAACAATCAAGAAGAACTATTTTAATAGCATAAATTTCTCTGAGATAGTGAAGATAAAAGAATGCTA
CCAGTTTACCCACTCCTTACTGTCTGAGTCAGGAAGAAGGTGGAACATCCTATTCTTCTTACCTCTATTACCTCATT
TAGAAAATTTTGACTATTTCTTCTTTTAGCTCCCTTTTACGGTCCACCATTTTGGGAAGATGACTTTCTCCCTGTCTT
CTTAGAATAAAAGGTCCAAAGGAAGAAAAGAACAGATATGCTCAGGAGTTACCTGTGATGACTTCTCTCCCTTGCCCT
CGGGCACTGGCCCACTCCTTTTCTATTCTCAGTTCTTTATCTATTTACTTTTCTATTCCCTGCTCCTTTCTCTATT
GTCCTGACTTGCATGACACTGGCAAGCTACAAAATGGGAACAATAGCAACTGCCTTAGAAGGCTGACAGGATTCACGGG
AAGCACCTGAGCAGAGTCTAATTCCTAATGAGTGGTCTTACATATTAGGATCATTTTCTTCTTCTTTGAGCTTCTGG
TTTCTTGGCCTCTGGCCTTAGCCTCTTCAACATATTCTACATAGGTAATCAGATTCACCTTCAAACAATGCTGTTTTG
ATCTTGTCTCTGGTGAAGAAAAAATAAAACCTTCAGTGGCTTTCTAAGACGATGAAATCAAATTCAAAGTCTCATCGT
GGTATTCAAGGTCTTCTGTAATCTCCCTCAGCCTGTGGCTCTTACTGATGCCCCATTCTTCTATTGCTCAGGGTGCTT
TGGGCTGTAGCCACTAGGTTGACGAGCATCCTCTTGACGAGCATCCTGATGATGCTGATAAAAGTGCTCAAGGACATTT
TGGGAATTTTTTCTTTAGTGTATCTAGGATAGTACTGTATACAATGCATATTTTATGTATAAATCTTATTTCTGTCT
ATGTTAGTTGCCTTGAAACAAAACCTGGGCCTTTGTATCTGTGCAGGCTACAAAGGCTTGACACTGCTACCCCTTAGAAA
GGCCTGCTTGCCAGGTTAGCCTTTGGTTGGTAACTGGGAACCTGAGCCCTTGGAGGGCTCTCAGTCAACAGTCAATTGAT
AAGTGTGGTTCACTGTGCCTAGACTTTTGTGCAACAACACAGTTTATGCTTGAACACCTGCTCCCTGTTTGGAGTCT
GGAATATTTTGATGTGCTAGGCAGAGGGTGCCTATGTGATCAGCTCCATGAAAAACCTTGGGCACCGAGTTTCTAAAA
GAAGCTTCTGTGGGCTGAAACGTCATATACAAGTTGCTACATTTTCTTGTGCTACTCTATGTGATCTCCCGTGGTAGG
CAGAAGCATAAGGAAATCTGCACGTAAATCATTCAGACNCTGCCTGTGGCTCTCCCTTATGATCTGGCTGTGTATCCT
TATTACATCACTAGAATAAATCTTAGCTTTAAGTACTGCCATACACTGAGTCCCATGGGTCTTCTAGTGATGTCCAAA
TGTAGGGGGCAGGGGTCTTGGGTACCCCTGACACAGTATCTTTCAAACAATTTCTTAGAGTAGGTTCTTAATAAAT
ATTTTGTGATTGAGCATCTCAAATCTCCCAACATTGTAGATGCTTAAACATTTAAACACTTTTGTCTTTTACCA
TTTTTGATTTTTTGGGCACCTTTAAGTTTTTTTTTATATGCCATCAGGCCAACATTTATAACATATGGTAGATCTGAGA
TGTTGTTTTACTGCCTACAAAGAATTGTTCTAGTTTTGCCTATTTATTTCTTTTACTTCCCGATATTTACAGACACTGT
TAGATCTTGCCAGAGCAGTTCTTTTGGGCTTTTGTGATACAGAATCACTGGAAAAGGTAGAGGGTTAGAAGGGCACAGG
TTTGAGGACTAGGTCTGGGAGATCAGTTTTGGTGGGAGGGTATGAAGGAGAGAGGGCGCATGACAGGGCTTGTGTGATGT
GACGTGACCACATGAAAGGAAGGGGCTTTGTGAATTGTTGCGGCAGACTGAGATTCCTTTTACAGCATTGTGAGAAATA
ACAGAAAGAGAATTTGTTATTTTGAATGACTTCCTGGCGGCTGAGTTGTTTTTCAATTTGATAACGACCCCTCTCTAATA
CCAAGTCATGGTTTGCAGTTTACGTCTTAGGGTTTTTACAGATTTTATTTGATGGTAGGAACCACATCATTTGTTCTAG
ATTTAGCATGTGAAATTCAGTAAATCTTGGTGACAATGACTGATTGCTTCATTCATAAACATTTATTGAATACTTG
CTGTGTGCCAAGCATCGTGTGTAGAGGATACAGAGATGAGCAAAAATAGGTCCCTACTCACATCATAGAGGGGAGGCCG
ACTCACATGCACACATTTCCAATCAGATATGGTGAATACTAGTGGTGAAGGTTAGGAGCCAGAGGACAGCATCTCAT
CCACTGGGACCTTGCTTCTTACGAGGCTTGTCTGCATTTTATGAGGCTTAAATTTACCAATGTAGTTATCTAGTGGTG
CTGATAAATCACTTATAATTAGTTTTTTTATTTGTCTAAGTATCAATTTTAGCTGCTGTTTTTTTAACTTTCTTTAGTTT
CTGCCTACATAATTTCTTGAATTAAGCATCAATAATGGACCTTTGCAAAAATAAGCACTTTCTAACACCAATGTGC
CATCCATCAGCAAAGGAGGCCATTAGCAAAGTCCAATTCATAGACTGCCTCTGATTTTGCATTTTAACTTGCTAATAT
TTAGCTCACAGTGTGATTTTCCCACGGTCACACACCTAGTTGGTGGTGAATTGCCGTAATTTTGCAATTCATATTATTT
TGCATGTATCTCATTTTATTTGTTATTCTGTGCCAATCTCTCTGATATTGAATGTAAGTTTATTGAGTGCAGATGTCTGT
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ATTAGGATTCTGTAGAGAAACAAAACCAATAGGAAATATATAGATACATAAGAGGAGATATATTGTGGAAATTGGCTCA
TGCAATTATGGAGGTTGAAAAGTCTCACAATATGTCTCTTGAAGCTGGAGAACCAGGAAAGCCAGTGGTATAATTGAG
TCTGAGTCCAAAGGTCTGAGAACCAGGGGAGCCAATGGCATAACTTCCAGTCTGATGCCAAAAGGCCTGAGAACTTCA
GGGAGAAATCTGAAGTCCCAAGAACTAGNAACTCCAATGTCAGAGCAGGAGAAGATGGATGTCCAGCTCAAGGAAAGA
GAGTTCACCTTCTTCTGCCTTATTGTTGTATCTAAACTGTCAATAAATTGGATGATGCTGGCTCACATTTGTGAGGGCA
GATTTTCTTTATTTAGTCTACTGATTCAAATGCTAATTTCCCCCAGAAACACCCTGAGAGACACATCCAGAAATAATAT
TTTACCAGCTATCTGAGCATCCCTAAGCCCAACCAAGTTGACACATAAAATTAATAATCACTGATGATAGTAATGAAAA
GAAAAAATTTGGATTTCTGCTTTTGGTCTTCATGAGATATTTTCTCTACTGTATCTCTCTTCTCATGAGATATTTTNCAC
TATATCACATGGATATATTTTCCACTTCGATATAGTAGAAAATATGCATGGAAAAATGCATCTATTATTTATTGAGCAC
ATATTATACCTTGAGGCACTCTGCTAGGCACTGTATTGAACAAAGATGATTAAGACAGTGTACCTTTCTTAAGGCACAGA
AACCATAAGATAAATATACAAATGACTGTAGTGCAAGGTAGAATATACAAATGCACAATAAGCAAATCTGAGAATGGG
GCTTTACACATTTGTGGGAGAATGAGAGAAAGCTTCATGGAGGAGGTAACCTTTTGACCTAAACCTTGAGAGCAGGAGCAC
CTGAGTTCAGAGGTTAATTAATAAATAAAGGAGAGGAGAGAGGGGAGGGTGGGCAAAGGGGAAAAACAGGAAATAT
AAATGTGAGTCATTTTTTGGAAAATGTGGGAAGTCTCTTTTAGGTGTGGTGTATGAAGTGTGTGTATGTTGGGGGAATA
GAGCAAAGTAAGGATTAAGAGTCAAGTTTCCAGGCTATGTTGCCTGGCCAAGCTCAGGGCTTTGCACTTGGTTCATCAG
CATTGATATGTTTTTTGTTTTGTTTTGTTTTTTTGGACAGCGTCTCGCTCCGTCCAGTCTGGAGTGCAGTAGTGCAAT
CTCGGCTCACTGCAACCTCTGCCTCCGGGTCAAGCAATTTCTCTGCTCAGCCTGCCAAGTAGCTGGGACTACAGGTG

Fig. 9.22

Fig. 9.23

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GATTGGATTCAAACCAGCACATTTAAAGGTCAACTGGTTGTTCTTAACTGTGCAGAATTTTAACTCTTTCATTATAAT
GGATTTTAAACCAATGGCCATTTTAGGAAGGAGAGAGACTGCATACTGTGTTTAACTTTCTGACCTTTTTGTTCTTAA
ATTTTTTGTAAAGTTCTCAGGGATTTCTATGGATGGAAAGGGTCTGAGAAATCAGTTCTCAGCTGGCATTGGAAG
GGTGGAAGTAGAACTGAGTTCTTTTAACTGCTAAATTTAGGTCAGGGCTTTAATACGACTGTGCACATCACATAAG
CAATGTCTAAATGTCAGGCTGAATGTTCTCTCTCTTTTAAATAAGTACTTCAAGTTCGTTAGTAATAGCTAGCCCTCTG
TCTCTTTCTTGCTTCTCCTCTCACCTTTGAGCAACTTTTAAAGTTCCTTTAACAGCTCTGTGGAGAAACCACCATGTAA
TGATCAATATTTAACCCAGNTGGTTGACGTTAGCCCCGTTTCTGGAATGGGTATTTCACTGGGCTTTGGGGTTTTGATG
ACGTTTTCTCCCCAGTGACTGTGGATGCATGGATTAAATTTGTTGGTCTGCAAAATACCATCTGTCTACCAGCCATTTT
GCTGTATTTCCCAAATAAGGTAGGAGACTGATGTGTCTCAAATAAGCAGGGTGAAATTTAGTCATCTGATGATAGTTT
CTTGGGTTGTGCATTAGTGTCTGATGAGCCCCCTAGCAACTTTCAACATTATGAAGACTAGTTTGGAGCCTCAAAGGC
AGTATCTTTCTTGAAGAAAAAATAACACCAGGCATTTCCACAAATACCATACACAAGTGGTCAGGTAAAGGCTGTG
GGAGTTCACGCATCTGTTTACCTTGAGTGTGTGTATGAACTCTTGTAGTTCTAACTAAAGGCAAATGGAACTGTATAG
AGTCCCATTTTGTCAAGCTGACAATTCAGTATCCTCTTCTTCTTCTGTTGCTTTATAGCCCACTCTTAGTTGTTNTCTT
TCTATATTTCTTACAATTTTGTAGCTGTGAGGCTGTGTATTCGTTGCTTAAAAATTTCTGTGCCTCATCTGCTTCACTTGC
AAAATGGGGAAATGTCACAGGATCCTTGGGGTGTGCTTTGCCAGCCAGAAACCTTTGTGGCTAATGGTGCCCTTTGCC
TGAGTTTTGCTCGGGCCCACTGGGCTGGTTCTGCCCACTCAGCCTGGGCAGGCTGTGCTGGGCTCACGCTACTGGCCCA
GATCTTATGCCTGCCAAGGGTGAGCCAGGCACAAAAGGCAAGGGGTATGTGAACAAGGGAGCTGCCCATTTGTGCACAG
CCAGTCATGCCAGCTGNAGCAGGGCAGGCAGCTCCAGATGCCGACACAAGTGCTAGCTCCCTGTGAGGCTGTGGCTGGA
CCAGGTGTACCACAGATGGCTTCCACTGCTGGCACCCGGGAATGTGGTGGTGCTTGGAGCTTGGAGACGCCAGGAAC
ACAGAGCCCCAAAGAGGGTGTCACATCCCTGGCTTGGGGAGCTCCCAGGTCTGGGCTCCCTGAAGGGCTGCAGTTCCTC
TCTTCTTCTCTCTTCTCTCCTCTTCTCACCTGCAACATGGCAAGCAAGGGGTGTGTTTCAGCCCTGTTTGTGTTACAG
CTCTTTCAGTCCCACCATTAGTGGGTCCCAAGTTCCTGTCTGTCTGGAAGAATGAGGTACAGGGACAACCTGGAG
GGTGAGCAAGGCAGAGAGGAACCTTTATTGAGCGACGGTACAGCTCCCAGGAGACCTGAAGTGGGTAGCTCCTCTCTGCA
GGCAGGTTGTCTGTCAAGCCTGCAGCTCTCAAGAGAGAGGAGACCCACAGTGGGTAGCTCCTCTCTGCAGGCAGGTCA
TCCCTTCAAATGTGCAGCTCTCAGGAGAGAGAAGACCTACAGTGGGTAGCTCCTCTCTGCAGGCCAGTTGTTCCAACCT
CTGCCCAGTCTGGCTGAGTCCAGGGAGTTTTATGGGCTTCAGAGAAGGGGAAGTGAGTGCTAATTGGTCTATGGACAGC
TGTGGGCAGCGTAAGTTCCTCACTCCAGTCTGTGGAAGTACAGCCTGGCCCCCAGGCTTCAGGCCATCCCAGGCCTAAC
GGTGGGGCTTCACGGAACTGCCCCCTTCTGTCCAGGAGTCCGTCTGCCTCCCGCTGCCATCAACCTGCTGCACTGGCA
CCCAGGCTGTTTATGCCAAGGGGCACCTGCAGCCCACTAGAGCCACTCTCAGCTCCCCCTTGGCCCCCTCCCATGCT
CCTTGGTGCACAGAGTCTGGAGGGGGCCGAGGCAACAGGGGGCTGGTGTGTCTAGTGGTACTCTGAGCACGGGCATACAC
AGCTGGGTCTGTGACAGTGCCCCAGCTTGCACCTAACCTTTGCTCTGAAATTGGAGTGGGAGCTAGGAGTGGGGAGAGGCT
GGGCAGTGGGAGCAGGCATTTCTGAGCCTGTGGGGGTAGGGGGGTTTCTTGGGCTTCTGAGAGTACAGGGATGCCTGGG
TCTGCACCCACAGCTGGGTGGCTGCAGCTGTGCCAGGAGGGCAGGACTCCCACCCCTCACACTCAGAAGTGGGTGGGG
CTTCTGCCTGTTCCAGCTCCCAACAGCTCCATGGAATGTGCAGCCCTGGCTGCGCCTTGCTGCTGCAGCTGGCGTCT
TCTCAGCAGCCACTCCAGATGGGCCACCACTGTCTATCAATTACATTACCTTACTGTTATTGTGAAAATTAAATGA
GTTAATCCATGTAAATGCTGGGAACAATGTCTGGCATAACATAGAAGCTCAATAAATATTAATTCTCATTTATTGTTGG
TATCATTTTTTGCTATAATTTGAATCTTTTACATATTGTTGTGACATATGGAGGTTTTTGACCTACAGGGTTTCTTCTGA
GTGGCATAGCTCAGGGTAAAGTGGCTCTCCTCACCTCTCAATACCCTACTGTCTTTTGCTAAATTCAGTTAAGGATG
GCTAGAGACATCATATGGGAAGGGTGGTCTGAACCAGAGAGATGGGAGAGCAAATGCTAAGGTGGTTCTTGAGGGT
AGGGATGGAGAGCCAGACAGAGACAGGTGTCCTCCAGACAGCTGTCTTTATCATTTCACTCACTGCCTCTGCAGCTCCT
GCTGCCTCCTTCAGAAAGCGAATGTATGTTGCCCTTAATGCCCTTCTGAGCTAAAGCCAGTAGTTCTCAAACATCCT
CTTTCAAAGACACTGAGCCCTGGTAACCTTCATTACATTGTTGTTCTCATATTTTTCTCAAGTTAACTCACTCATAACC
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TTAATATTCACAAAACAGGCTTAATCTGAGCACAAAGTGAAATGGTATAGCTGCGAAATACCTGGCAGTGGTAGGTACTA
GGTGTATAACATGATGACACTTAGAATTTATAGACACTTAGAGCATAGTTTTTCAAATATTTTGGCTGTAATCAATAA
GAGATAGATTTACATCACAGTCTATACAGATGTACATTTATTTATATGTATGTATGTAGGCATAGCTTAAGCAAAAGT
GTCATGAAACAATGTTCAACCTATTGCACACAATGTACTTTCATGATTTGTATTCTCTTCTTTTTTTATTGAAAATGAGT
CTGGTTTGTAAACCCAGTAAACACATAACCTACAGTTTGAAAAACCTGATCTAGAGGAACATCCAGTGGGTGCCTCTTG
AGTTCTTACGCACTATCCACATATCATCCGTATCATAATAATGGGTGTGGCCATCTCAGTTTTATTTTGAGGCACTG
CCTCTTCTCATCCTGACAGAAAAAAGAAAAAAGAAAAAGAAAGAAAGCTCAGATCAAGAACTAACACC
TTATCTTAAACATATAATCTCTCTTACCTTATGCTCTTTCCTAATAACTATTCTTTTTTCGATTGTCCAAAAGAAAC
CACTCTAATAACTTATAACACCATGTATTAGAAATTTAGAAACCATTTTTGAAAGACAGTAAAGATGTAAATTAATGGG
GGGACTAAATTATGCATGGATAGAAAGACTATTTATTATAATGACATCAGTTCTCCCATAAATCAATGATATTTCAAT
AAAAGGGTCAACAGTCTTTATTTTATAATGGAGACTCACAATCTTATTCCAAAATTTATATGGAAATGCAAGTAGCCAA
GAAGAGGCAATATAGCCCTAAAGTAAAGAACACGGAGGAAAAAATTAACCTATCTGATATCAATACTTACTATAAAG
TACAATACTTAAGTGTGGTACTGACACAAAGAAAAATGGTTCAGTGCAGTGATCATATACATCTTTTGTAGATGTATT
TGTATGTATTTGATATTTTGTATGCTGTGCAAATTTTATTGTTTTGTAAATTTTACCCTGTGCAAGTATCAGAAATA
CAGTTGATTTTTCTATCCAGCAAACCTTGCTAACTTTGTGGTTAATGGAAACACTTCAGCTATAGATTCTTTTCAATAA
CACAATCACATTGTCTAAGAATATTTTTATTGCTTATTCTTTTCAATCCTATGCCTTTTCTTGTGTTTTCCCCCTT
GCTTTATTGCTTTGGCTGAGATTTCCAGTGTCAAATAGAAATGTTGATGGTAAGCATTTTTTTCTCATTCCTGATTTCA
GAAGAAAAGCTTGTAGCATTTCAATATTGAGTTATTTTTTTTTGTGGTGTAAACAATTTTTTTTTGTAAACACCTTTTATT

Fig. 9.24

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ACATTTAAAGAGTTTTCTTCTGTTTCTAGTTTGCTAAGAGTTTTCTTGTTAGACGTTAAATTTTATCAAATGGCATTTC
TGAATCTGTTGATATAATAGAAATTTCTCCCTCATTCTGTTAATCTGGTAACTATATTAAATTGATATTTTGAATGTT
AAACCAACTGTATTAGTCTCCAACCTGTAGTAACTAACTGAATTTAGTTACACTGTGTTTCATTACTGGATTTCAGGT
GGTTAACATTTAAGATTTTGCACCTGTCACATGTAAGAGAGTTCTGGCTTGACTTTTTCTTTTCTTTTAATAACTTTGTTG
GGTTTTGGTGTAAAGGATATGCTGACTTCATAATAACAAGTTGGGAGTGGTTCCTGTTTTCTCTTCTCTGTAGAGTTTA
TGTAAGATTACTGTTAATTCTTCCTTATCTGTTTGGCAGAATTCAACAGTGAATCGTCCATTTTCATATAAAACAAGTTTA
TTTGCTTAAGGTTACTCAGAATATAATTTTAATATATCTGAAGGATCTTCAGTTAATGTCCCTTTTTCATTTTCAGATAT
TGGATATTTGTACCTTTTGTATTTCCTCCTTTATCAGTCAGTCTCACTAGGGGCTTATCAGTTTTATTAGTCATTTCA
AAGAACCATCTGTTGGCTTTTTATTGATATTCTTTATTGTATGTTTTTAAATTTTTCATTGATTTCTATTCTTATATTA
TCATTTTCATTTCTTCTTTTCTTTGGACTTAATTTGCTGTTCTTATATAAATTATAAATTATATAAATTATTGAAATATAT
GTTAAGATAATTGATTTTTAGGCTTTATTCTCACTTAATATTTACATTTAAGACTATAACATCTCCCTCAAAGGCTGGAT
TTAGCTATATGGCACAAATTTCTACTGTAGGATTTTCACTTTTATTTCAGTTCAAAATGTTTTCCAATATCTGTTTTGA
TTTCTTCTTTGAGCTACATGTTCTTTATAAATATATTACCTAATTTCTAAATATATGGGGATTTTCCAGTTGTCTTTTT
GTTATTGATATCTAGTTTAATTCACCTTTGGTCACTTTAAGAGAATTTGAATTTCTGCCATTACTAGGGCTGTGTTCTAT
ATATGTTGAAGTTTTAAATCATGTTGTTCCCATTTTCTCTATAACTGGTGACTATTTTCCCACTTATTAGTCTGTTAC
AAGAGGTGTGTTAAATGTCTTATGATTTGTGGGATTATCAGTTTTTGTAATTCTGTTTCATTTTTGCTTTAAATAATTT
AAGCCTCTGTTATTAGGAGCATATACTTTTAAATTTATTATATCTTCCTAGTAAATTCAGCCATGTATTTCCCTCCTCTT
TATCTATTAATGTTTTGTCTTAAATCTACTTTGTATGGTGTGAACATGGCTACTTCAGCTTTATTCTGGGGTAGTGCC
TGCATAGCGACCTCTTCTATTCTTTTACCTTCAATCTTATATAATTAAAGAAATGTCTCTTGTAAGAAACATATAGCT
TTGTTTAATACAGCCTGGCAAATTCGTCTTTTGAATAAAATCTTAATGTTTGTGATTTATGTAATTATTATATTTGTG
TTTTGTTTAAATCTATCATTATTATTGCTCTCCATTTGTCTCATCTTTTGTTCCTTTTAAATCTTTTTTACT
TTCTTTGTCTTGAGGTTTAGAAAATTTATTATTATTTCCACTTTTCTGCTCTGTTGACTTCCACCACTCCCAACTTA
TACACTTCTGTTTATGTGTTTTCACTTGGCATCAATATTAACCCCTCAGGACATTATTATCAAAGACAGTCAATATCT
ATTTAGATATACCCACATAGTTAGCCCCCTTTTTGACTCCTCGTTTTTCTGTATCTCAGTTTCCAGCTGCGATCATT
TCTGTCTGCCTGAAGAGTACCTGCAGTGCTTCCTTGAGTACGTGAAAGCTGCTGAGGAATCTCTCAGCTTTTATTTTTC
TGAAAATGTCTTTATTTTAGCTTTGTTTCTGAAATACATTTTCACTTGGTATAGAATTCTAGGTTGAATTTCTTTTCC
TTAGGCATTTGAAGATGTTTCATTGTTTTCTGCTTCCATTGCCATCCAAAGTCTTCTAAAACTTCCCTAGGTAGGTAGG
TAGAAATATTTAACTCAAAGGATGAATAAAAATGTATCCACAAACCCATACTTCTTTTTTAAATGGGATTTAAAGTTTA
TAGATATTTAGTATAAAGTATTTTACATCTGCAGATTGAATGCAGATGATCAAAGGAATCAAGTATTTGATGATTCAA
AATAGAGACCTTTGTTTTACATATAGACTAAGGGTTGGTCCAGGACTATCAAAACAATCTAGGAAGTATTTTCTAAC
TCTTGAAGAGAGAGAGAGGGAGCATAAAATGTACATAAACCTAAGTTAAAGAAGTATGTAAAGTATGTTAAATAA
TGCAAAAAGCATATATGCATATATTTGCTTGAACCTTGATTTCCACTGACTTGGAGTAGTTTCATTCTCTAAGAATCTCA
TGTCATATTATTTTATATCTTTCTCATTGTGAAGTCATTCAAGAGATCCTGCCTTGATGTGTTTTCCAGATAATTTA
CACTTTTATTTTACATAGATGTTGATTAGCTGTGTTTCATTGAATATTCTCAGTTTTGGGTATCAGTTTTTCAGCAAAA
CAACTAAATGTGACACCTTCTACTGAGCATATTGGGTCTATACGTGTGCATTTGACTTACGACTTATATTTTCACCTA
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AAAATATTTTAAATTTTGTATTTTAAATTTTGTATTTTAAATATTTTCTGTACTAAATGCACTACAATATAA
TGTGACAATTATGAATATGGATTTTAGATTAAAGACAAACCTGGGCTGGAATAGTAGCTCTGTTTCTTACTAGTTGTGTA
TCCTTGGAAAAACAACCTCCAACCTTCTAAGCATTAGTTTCTTATCTGTAACACAGGGTCCATAATTTCTACCTTACAA
TGCTGTTTTTAAAGAATAAACGAAGTGGGAAATGAGTTAGTATCATATTCATATATGGCAGCCATTATTATTATTATT
ATTATTATTATTAAATTTCTATAGTATGTTATTGCCCTAAGTTTGTTCATAGAATAATGTATTGGCAAATAATATTCAG
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AACATGGGAGTCAACAGCNTATAATAAATCTGTATTCTTAATTTAACAAGCATTTATTGAGTCTCTAACAACAAGCTT
AGCAGTGTTTAAACACCATGGCAGGGCTGGGCATGGTGGCTTATGCCTGTAATCCCTCACTTTGGGAGGCCAAGGTGG
GAGGATCACTTGAGGTCAGGAGTTCGAGACCAGTGTGGTCAACATAGCAAAGCATCGTCTCTACTGAAAATACAAAAAT
TATCTCGGCATGGTGGCAAGCACCTATAATCCAGCTACTTGGGAGGTTGAGGCAGGAGAATTGCTTGGATCAGGGAGG
TGGAGGTTGCAGTAAGAACTCGCCACTGCACTCCACTCCAGCCTGGGCGACAGTGCAAGACTGTCTCAAAGAAACAAAA
CAAAACAAAACAAAACAAAAAACTACCACGGCAGGAGGAATTTCAAGCATGTGAAAGCTGTTACCAAGGATAATTGTG
CCTCCATCACAGGTGTCTGCCTCTCCCCATCTCTGCTGTGCTGGAACCCACACAGTATCACTTGTCTGGGTTTTCT
ACAGTAGCTGCCTTACCCTTAACTCCCATTATTATTGTCTCTCCAATCCATCCTTTATACTTTTTTCCAGAAATTATCTTT
CTAAACAAAATCATGCCATCATCATTCACACATTCATAGACAGCTGTTGTCTGTGGAAACACTTCAGCCTCATTAAG
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GGACTAATTGCAAATCTCCATTAGTACCGTGTCTTCCAGCTGTTATTTTAGCTAGACATGTTTTTTTAGCCATTTCT
TGCTCTCAAATTTGTAATCTTCTTAAAGATATAAGAGAAACACCATCTTATCTATAGTCTCCCTAGATGTAGAGGAT
GAATTCGAGTGTTAAAGCAAGAGGTGGGCGTTGTAGTGTCTAAGGTAGGGAGACTAGGTATAGAAAGAAACCTTGAG
ATGGAAACCAAGGACAGAGAACTTTTGCAGAATGCTGTCTCTGTGGGCTGCCTGCCCTGTCTTTCAGCTGGAAGTGGT
TCTTCCCTTCTGTGTTCTCTTTCTCTGATGGGCTGCTGAGAATTATTGCATGTAGGAAGCCAGAGAATGTCTCACTGTT
CTCCAGCAGCTGCTGCTTAGGGCTCTCTTACTCCACTCTTTTTTGTATTCCCTGGTCTCCTGCAGAGCCATTTATTGTC
TGGACCTTCCCTATACGTTGTCTTCCCTTAGCTCAAGGCCTGGCCTCTCCTTATCTCTCCTCAGAGTTTGAATTTCTGA
TGGGATCTGTTGTGCTGGAAGTAGCTGTGATGTTTCTCTTCTTCTGCTTACCTATGGGTTGGTTCCTCAGCCCTTAT
ACTGGTCCCCTGGCACCAGCCCTGACCCGATTAATTCCTTTTCATCGTCCCTCCATACCCAAAGGTCCTGTCTTGGACC

Fig. 9.25

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AGGGAGGCTCACTGGGCCAAGTATTCATCACAAATGTTAATAAATATAACCTAATTTTAAAAGAAATATCTCCAAAGAG
TGTAGCCTTCTAGGAGACTGAGTACAAAAAAGAAAGGGGTGGAGCAGGACAGAGTATGAAAGAAGACTGCAAGAAAA
GGTCAGGTACAACCTGGGAGAGAAAAATGCAGAAGCTGTGGGCATGCAAGGCCAGAAGTGTAGCCAAGAAGCAGNAGGTG
AAGTCAAAGGTGGATGAAGGGAAAGAAAGATGAAATGAGAGAAAAATCCTAGGAGTCTTAGCATTGGAGGGGGAAGCTC
AGGTGCGAGAAATGATCTAATAATAGTTGAATGGAGAGAAAAATCAATGTATGGTCAATCTTCATTATCACAGATTATGT
GTTTGCAAATCCACCTACTTGCTAAATTTATCTGTAATCCCAAAGCAATCCTTGCGGCGCTTCTGCAGTCATTTGTG
GACGAGCGTGAAGCAGTGAAAAATTTAAGCAGTGCCACATGTGTATTTCCAGCTGAGGGTGAACAAGGGATGCTCAGCC
ATCGTGTTTCAGCCCTCATGCTGTAAGCGAGGGTCTTTCCATGATACGTTTAATGCTGTGTTTTTGAATTCTTGTTGTT
TTTCACTGGTGATTTTGTGTCATGTGAAATGGCTTCCAAGCATAGTGCTGAAGTGCTCTCCAGTGCTCCTAAGCACAAGAA
GGCTGTAATAAGAAGAAATCTGGCCAGGTGCCCTGGTGCATGAGACCAGCCTGGCCAACATGGTGCAACCCCATCTCTA
CTAAAAACATAAAAAATCAGCTGGGCCTGGTGGTGTGCACCTGTAATCCTAGCTACTCGGGAGGCTGAGGCATGAGAATA
GCTTGAACCCGAGAGGCGGAGGTTGCAGTGAACCGAGATCATGCCACTGCACTCCAGCCTGGATGACAGAGCAAGACTG
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AACTTTGTTCAAACATGAGTTATAGAGCTGCTGGCAGGGTGTGTTAGTGTTAATGAATTAACAATATATATTAATAAAG
CTATCTTTAATAGAAACACACAAAAAACAAGTTATGTATTGATTAGGTGATGAAAATATTGTGACCAGAACTTGCAG
AAACCTGATCTTACATTTCCCTGGGAGCAATGGTTCAGTATTTGCTAATTCAGTGTTCTCAGTGACTTTATAGAATGT
AACTGCCACAAATAACAAGAATCAACTCTGTAAACATTTTTGTTGGTCAATGGATATGCATGTTTAGACATTTTGTAGA
ATTATAGTCAACCCAAGTGAAATGGAAGTCTTTTATTTGACTTGGGAAGTTGCTCTATTCTCTTCTGTTTCAAACAAA
ACTCTAAATAAAATCCTCTTTTGTCTGGTTTATTTTAGCAGATGTTTACTGGTGTGTTACTGGCAAATGTGTTGAGTAA
GTTAGGTAAAGTTGACCTAGACACAGTTTTTGTCTCCCAAGGAATTCTCAATTGATCATGGAAGACAGGAGATGTGTGTA
ATCAACAGTATTGACATTAATTTGTATTCTGGAACAGGTATCGACTAGAGTCAAAGAAGGTTTTGGCTGGTCTGAAC
AACAAGGAACCCAACGTAGGTTACAAAATAACCACATATTGTCCATAGAGTAGGTGTCTACAACATGCTAGTGAAGGT
TGACATTATCTTAGAGCTTCGGGGATTTATAGATGGAAGGGATCCTACAGGCCAGTATGTGACAAGGACCTGAGCATG
TTTAGATGTAGAGCTAGAACTGAAAACCAAGCCTCATGGCTGAACTGAATCCTTTCCCACTAAACCATAACAGGCTCTA
TGCGCTCGAACTAGTTATTGGAGTCCCGGCTTAAATGAGCTCATGTTCCAGAAATTCCACTTAGGTCAGATGTTTATA
AGTCAGTCTGAGGTTGTTCCAGGATGAGCTCCAAGGGGTCTGTTTTGCTGTATATGAATATTTTATAGAAGGAAGCTC
ATAAATTTTCATGTGATTCTCAAAGGAGTTTCATGATTCCAAAAGTCTAAGAACTTCTGTGCTTATTCTGAGGTTAACT
GTGCACCACAAATACTAAATTTCTATTTGGGGTGGTGTCTTGACATTGTATTTTGTATCCCTATCAATCTAGCAGAAGAT
TTTTTTTTTTTAAATAAAGAAAATGCTCACTCTTGGCTAGGATGTGGTGAACAAAACCTTCTATATTGTGAGTGGCAGT
ATAAATTGGTATGAATTTTCTAAAAAGACATTTTCCACTTTTTGCAACCAGATACCAAATTTGTCTTTCAATAATATATT
CTAAGAAAGCAATAAGAAAAAGGAAGTAATTAGAAATTATGTATAAAGATATTTATAGCATTTTTGTTTATAATATGAA
AACTAGGAATCAAAGTGTCTAAGAATTACAGTTTTTAAATATCACTTAGGCCGGGTGCGGTGGCTCAAGCCTGTAATCC
CAGCACTTTGGAAGGCTGAGGCAGGTGGATCGCCTGAGGTCAGGAGTTCGAGACCAGCCTGGCCAACATGGCAAAACCC
TGCTCTCTACTAAAAATACAAAATCAGCTAGGCATAGTGGCGGGTGCCTGTAGTCCAGCTACTCAGGAGGCTGAGGCA
GGAGAATAGCTTCAACATGGGAGGCAGAGGTTGAGCCGAGATTGTGCCACTGCACTCCAGCCTGGGCGACAGGGTGAAA
CTCCATGTAAAAAATAAATACTTAATATTATGAAAACTGTTTATGACAAGACATGATGTGAAAATAATCATGGCA
TATATAGCATAATCACAATTTGTTTATGTGTGAACAGAATAACATGTTAACAGAGGTTATCTTTTGCTAATNGGACTAG
AGGTGTTTCTTTTCTTTTATTTTTCTTTTGTATTTTTCAAATTTGTCTAAAAGACTATATATTTTTTATAATCAGAA
GAAATATTGCAACCATATTTTGTCCANTTGAACATGATTTACTTACGGGAAGTCCTATCCTTGTTTCTTAGTTTCTA
CATTTTAAATGTCTCATCATTTTCTTTGGCTGAAAGTTAAATGCAATCTCAGTTATTTTACGTTAATTCAATTACAA
GCCCTTCTCTTCCACAGTATCATTTTCTTAAACCTTCATACTCAGCCTCCTTTGCAACTCTGGCCTCTCTTGCTTT
CCATTCTTTTTTGTCTTTCTCTGATTAAAAAATACTAAATTTCTGTCTGCTTTTCTTTGACAGCATTAGCCTTTTT
TCTCCTCTGCTCAGAAATATAATTTTGTCTATTGAGTTTTTCAAATATTTACCCTGTTCCTTTGTTTCTTTATCCTGG
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TAATTAAGAAAGTATGTATTCTTTTTTGGGATTTCCCAAAGTATTTTCCCATTTATATCAGGAACACATGTGTATATGTGCA
TACATAGGGTATGCATGAGAATGAGTGTATTCACGTGTGCATGTATGCATTCATGTTTGTGTGCATGTTGGTGCATACA
TGTGCAAGTTGGTGTGCACTTGTGAGTGAATACATGTTCCAGTTCTTCCAGAACAGGAAGTGTGTCTTTCTTATATGC
CCTACGCAAGTAGACACTGCAATGATGCTGACTGCCTGTTTGCCTGTCTCTGTATAGCCACCTTTTCCAGAGCTTGCCATC
TGGATCTCAGACAGTATGCAGAGGAGAGGAATGTTCTAATCCACCCTGGTACCAACAGGCTGGCATCTGTACTTTGAAA
GCTTTGATGAAGAAGATGCAAAATGTGTTGGCTGTGTGGGCTTTTTCAGCTGCTTCTGACTGGGCTAGAGACAGGCAGC
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ATATGTATGAATGCAGACTGAATTAATAATGGCACCCCTAAGGGAGAAGCCAAGGACCAAAATCAGGAGACCCTCTAACTGA
GTCAGAAATCAAGGCCAGGATTTATAGGATGAACATTTTTAAGTGTACATTAGAATGTAAAGTAGGTATTTGTAATTGA
AGTACTATGCTTATTTTATTTCCATAAACCTAAGGTCTGGCTTCCAAGTGGCCTCAGGGTATATCCGTCAGCACTGAGC
CTTGTAATTTGCTGCTAAATCAAACCTATGATGAAAAGAACCAGGCAATAACACTGTCTTGCAAGTGTGCAAGTCCC
GTGGCTGTATTATAGAGTGCTCAGCTCAGTGTTTACAGTGGAGAAGAAGTACGTTGATATGCTTTCTGGGTTTGAGA
GCTCTTCTTCTTCTCATCCAGAAGACAGAGCTGGGTAGAGAAAAGGATAACCAGCAAGACAAAGGCTTTATCCCAATATT
ATATGTTTAAACTCAAAGAGATATGGTGAAAATGGGACTTTATTAATTTGCTGATTGGAAACAAACATGCAAAATTG
TACAAGATTCCATAATGTGCCAAAATGGCAAGGAGAGAGCCCTGACGAGCATGGAGAATAGTGAGTAATGCTTATGTCC
TAACAGTGCCCTTGCCTTCATTGACCTTTGGAAATATTGTGAAAAGTCAACCTTTTCTATTCTGTCTGTATAGTCTT
ACAGTGTTTCTCTCTTTTTCTCCCTCTAGGGAGTTATCCAAAGCCCCCTTCACAAGAAATGCTAGCTTTCAGGAATAA

Fig. 9.26

Fig. 9.27

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CTTTTGGGTTCAAGTGATCCTTTCTCCTCAGCCTCTCCAGTAGCTGGGACCACAGGTATGGCCACCACACCCAGCTAAT
TTTTTTATTTTTCTAGAGATGGTGTCTCACTATGTTGCCTAGGCTGATCTCAAACCTCCTGGGCTCAAGTTATCCTGTTT
CAATCTCCCTAAGTGCTGGGATTAGAGGTATGAGCCACCACACCCAGCCTGGGTTTTAGTTTTCTTTTTTGATCATATTTG
GAGAAAAAAGAAAAGAAGCAAATTTGCTAATCAAAATTGAGTGAATGTTTCAAGAAACAAAACCAATTAAATGTAGTC
CAACCTTGAGAATCAGCAAGTAAAAACCTCTGAAGCCAAGAGGGCTAACAGGAAAACCTGAAGGAGAGGAGAGAAATATA
CCATAAGCATTTCCTCAACCTCTGCTTCAAATTCTTGATTTTATAAATGGTCTGGGCCAGCAGTGCAACTCTCCCTGTCT
CGCTCTTTGTTGTTTATTTTTTTGGAGTCTTTAATGACCAATCGGGTTGCTGAAATTTACTTGAAGAGTTTATTCTTGAT
TCTGAGGTATAAGATTTTCTGTGTGAAGATGGCTGGGAATCCACACAACCTCTCCTAAAATAAGCACCAAGTTTTGGAAA
GAATTTTCTCTTGTTGGAACATCCATTTCCAAAGAGAGCCATCAGCATGAGATTGCTTTTTTTTTCTTTTTTAACCGCCC
CCAACCCAAGGAGAAAGCTGAGGGGAGTCAGTCTGTGAAGCAGCCAGGAATCAACTTTTCATGACAGTTTTTAACATTCAA
GGCAGACTTTTCCCTTGGTCTTTTGGGAAGAACTGCTGAAATCTCAATGAAAGTTCATGAAATTTCTGTAGAGGAATCAGC
TGAGTCTAGAAATAAGCTGTTCCCTTATTGTTTCCTCTGTTTCCTTATATATAAAGAGTGGGTAAAAAGAAGCACCTTGG
AAAGTCATTGTGAGATTAAATGGCATAATGCATGCACATCTCCTGGCACAACTAATCACCACTGTTGTGTTCTAAGCC
AGTTTCCTAAATACACAGATCCAGTTGTTTTTTAAGTCTCAGTGTGAAGTGTGTTTATTCTGTTTGGCCCATGTGGATATC
TAAAGGGTTCACAAAAATGCTTTCAAATTGATAAAATATGGAGGAAGCCGCTTTTCTAGTCTTCATCAAGGATCTGCGT
TGGGTGGAATAGAATCTTTGTTTGGGTTTGGTAAGAGTTTCATGATGCAGGATTTTATAGAGAGGGAGCTAATGCCATT
GGGTGGCTGATCACCTAGGTGGAATGAATGTGGGGCCACTCATCCATGCATTTTTCTGCAGGTGAGATGGGATAGGTAG
CCCTTGACAGTTCTTCTGTGAGAGATCTTGGTCACATTTGAGTCTCTGCTGGGCTAAGGCAGAAGCCCTTATAGGAGG
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AGCCAGTCAGATCCATGTCCAACCTTAGCAGCTAGGATGCTGGCCATCCCTGTGGCCAGGCCCCATTGTACAAGTTC
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TCTCTAACCTGAGTCACAGCACAATCTTTGTGGTCTGACTCTGTCTGGGGCATGGAATAATTGTGTGTGTGTGTGTGT
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GTTATCCAACTTTTCTGTGCAGAAGAACAAGATTTAAGTTTGAATGGATCATTGTTTCTTTTCGGCTAAACAACCTGCT
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TGTTCCCTTCTTGGAACATATGTTATGGAACCTTGACTAATCGCCCTGGCAATATGTTGCCAAAGTTTTCTTGCAATAGA
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TTGCAATTAATTTTGGAGTCAGATTTTGATAGAAAAGTAGAAAAGAGGATAGGACTAATTGACTTCTGAAAGAATTACAN
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AAGACAAAAAAGAAATTGCAAGCGGCCAATACATAAGAGAAGATGCCTAACACCTAACCCCCCAATTAGCATTGTGATTA
TAAAGATAACAATGCTATCATTTCCCCACATCATTTTGGCAGAAATGTGAATGGATGGTTAAACTCGGTGTGCTGTGA
GTTTGATGAAATGGATGTGCTCATACTACCTGATACAACTTTTGGAGGGCAGGTGGCTGATACTTACTGAAATGT
GAAATGTATAATCTTCTTGCCCAGAAATCTGGGTCTAGGACTTATCCTAAGAAGACAACCTATACAAATGTAAAAATAC
ATTTATGTAGTATGTTTGCAAAAGCATATTCATGGGGAAAATGAGAAATAACTTTAATGTTTATCAATAGAGAACTGGT
ATAGTAAATTATGATAAATACCTACTGTGAAATCCTGCATAGTCAATAAAATGATGGTGTAAAGCTTCATATATTAATG
TGAAAAATTATTTAATGGTACAAAACAGGTTATGAAACACAACAATCACATTTCTATACAGTCATATTACTAGAAATAT
GGGAAGTTGTTAATTTAAATCTGAACAGTTAGAATTTTCAGGCTACCTTTACTTCTTTGCTATTTAACTGTTTCTCTC
CCACCTATCAACTCTATGTCCTTAGGGAACTTATTTAACCTTTCTCACTCTTGGTGAAAATGGAGATAATACCAAACCT
CTTAGAGGTTGTTGTGAGAATCAAGCAATAAAATACATGTCAAGCACCTAGTAAAAAATGTATTTTGATTTTTTTTTTGT
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TACTCAGCTGCAACCATCTGCAGAGGGCCACATTGGCAGCAAAAGGAAATAGCCAAGACTCACAGGAGCACAAACAGCC
AGTCAGGTTTGGGATTCATTTCTCTGATACTGTGCCAAAGTTGGTGGACTCAGCCATTCTGAAACTCACTGTTCACTTG
GGCCTGCTTTGCACAACAAAACAAATAGCCCTGTTTTGCTTTCCATGTCTGAGGCAGAAAAATAAAAAATGGCAATATT
TTGAGTGTTTTTTACATTGGGAATTTCTGGTCCATAGATAATTATTTTCAGATACCCCTGTGTAATCTTTTCAGAGCTGTT
GAAAACGTGGAATAGGGGATAATTAGGGCTCAAGAGATTTCAATTAATGGTTAAAGGATCATTAGAAGGACACGTACTG
TATCCCTCTTCTTATTTCTCTTCTGCACTAAAACAACCTCCTGGCTTGCTACCATACTTCCGCACCCATGTGCAATTTT
CACTTAGCAAAAGGTCAAGTCTAAGTAGAATCTATGAAGTTGGGATTGATTCTGATTGGTAATAATGCTAGCTAGCAAG
CATTGAAAATTTGTACCAGGTTTTCTGTTAAGTAGTCTGCATGCACTATCTCAATGATTCTTAAGGAGTGGTCCATAGA
TCACTGGCTTTACAAACATTGACGCATGGTGGTGGGTGTGGGCAATGAAAAGGCAGATTCTAGATTCTACTCTAGATC

Fig. 9.28

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CATGGGATCAGCATCCTGTAGTGGGAGCCAGGGATCTACCATTTTAGCAGGTGCCTTGGGTCATCATCCTTATGCACAT
GACATTTCGAGAGCCACTGTGTTATTTCTAGTGCAACTGTGACTCAGTGTAATCTGCTTTCTTTCCACTCCTCTGATTT
CTGTGTGTAAATTTTCATGACAAAAGTTAGACAAAATACCATGGACTGAAGTTAGATATAGAGCAACTCATGTCTCTTT
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GGAAGGTTAAAGAAGTGAGAACTCTTTCCTTGCTTAGTGGTGGCCTGTGCTCTGGAGGTAGAGAGGAAGACACAGGGGC
CATGGAGGTGGGACTTACAGGTGTGGTACAGGTTGTATAATCATGATGGTGACCCTGCATGGAGAAGGGAAAGCTTGAC
CTTTTCCTTGGCTCAGGGCTATGCAAATAGGGCTGGCCCATTTCTGAGAGGGGAAGGTAGCTGATATGCAGGTGTGATT
CCCTAGGGCCCAGAAAGACCTGCCTGCCTTCTGCCTTTGTTGTCATGCTTCATGAGGCCTGAAATGTTCACTGGTGGTT
ATTTCTCTGGTTCTCAGCTTTAGCTCTAGAAAGCCCCATTGAAATGACATGCCACTTGGAAGATAGCAGGACCATTTA
CTTATATTTTCACAGACTGCTTTAATCTGCTACTCTGGGGATGGCACACAAAAGATTTAGTTGATGGGAGGGAGGTTA
GGCCCCCTTGTGGGCCACAAAGAAGAATCTGGGGAGCCAGGGAGACTATTCTGGGGGTAGAAGTGAGGAGGAAAGGCT
CTAAAGACAACTGGACCTATGTCACAGTTCCGAGGGCTGTCCTCCAGAATAACATTGTTTTCTGAAATTCTCCAGCCC
CTTCCATCATTTTCTCTCAGTGAATGTCCAGTGCTGAATAACTGCCCCCTTAGTACAGGACTGGACTGCAGCAGTGC
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GGGAAGAGGGAGATAAFTACCCACCTCTGAGTCTGAGATCCCCCTACCCCCAGTGCATTCTTCTGCCAGGAGCAGGAT
CATTAGAATAAGTGAGACTTAGCTTAGCTCATTCAACTCCTTGCCAGGCCCCCATGCAAAGCTTCAAACCTTCAAATAT
ACGATCATTTCTAGAGACTATGTTGGGATCCCCCTGCATTTCTTTATCCATTTGTTGAGAAGGACCAGAGATGATGCCTA
ACTTTAGCCAAGAGTTTGTCTGTGGGTCCGATAAGCCTCACATTTGATTTTATTGGAATTTTAATCTTGCAGAAAGTAC
CCTATGTTTCAATTTTAAAGCATTTTTTTTTTGTTAAACTGAAAACATCCCTCAATTTTTTCCAATGTTTCGTTTGTCTCTA
GTGCTAATTTATGTGGTAAGTGGTACTGGTTCATTAATTTACTCAGGAGGGTGTAAACCCTTTCATGGAGAAGGGGTG
GCTAGGAAGGGGTTTCTAGTGAAGGAATTCATTTTTCTTGGTCTTCATGGTGGGTGCTGGGGAAGGGTGTGCAGTCAGA
GGGGCTGCATTTGCAGAAACATGGCTCTGTGTGTGAGCCTGGGTTCTCAGGAATAGGAAGCAGGTAGGTGCAGTGGGTG
GGTATTCATGTGACAGTGGAGGAAGATAAGGACAAAGATCATGCATGAGTAGTGAATGAATGAATGAATGAATGA
ATGAATGAATGAATGATTGTGTATAGAAGTCCATACTCCTTCTCTGGGGTGGCTATTCTTCCACCAAGCAGACTCTGTT
CTTCTTCTGCTGCTTCTTAGTGAGCACGTCTAAGTCTCAGTTGATATTCTCTTGCTCTGAGAAATTTCTCATCAATGG
GAGGAGAAAATATAATTCGCTTCAACATAGCTCATATTTAAGCTGAGAATTCAGCATGAATTCAGACATGGTTCATGT
ATTTTGGATAATCAGCATAGCTGTTTCATGATCGGTAACCCTCTTTTTTCTCCTCCTTCAAATCGTTTTTGGTTAGGTTA
CCATGACTGAAGATTAATGACTTCCATTATTTTTTTTTCTCATGCAGGAATGTTTAAACTAGTCTAAACTTTGTACCA
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CGTGTGTGATGTTCCCTGCCCTGTGTCCAAGTGTCTCATTTGTTTCAAGTCCACCTATGAGTGAGAACATGCGGTGTTT
GGTTTTCTGTCTTGTGATAGTTTGCTGAGAATGATGGTTTCCAGCTTCATCCATCTCCCTGCAAAGGACATGATCTCA
TCCTTTTTTATGGCTGCATAGTATTCCATGGAATTCACCTTCATTTTCAATATTGTGTATATTTATTGCAGTGAGCTTA
ATCCTAATAAAAACAATTTTATTTCTATTTATTTATGTTTTTCAATTTTTTCTTTGAGACAGGATCTCACTCTGTCA
TGAGGCTGGAGTGCAATAGCACAAACCATAGCTCACTGCAGCCTCCAACTCCTGGACTTAAGTGATCCTCCCACCTTGGC
CTCCCAAGTAGCTGGGACTACAGGTGTGTGCTACCATGCCAGCTATTTATTTATTTGTTTCTTTTTTTGCAGAGATG
GGGTCTTGCTGTGTTGCTCAGGGTGGTCTCAAACCTCCTGGATTCAAGTGATCCTTGGCCTTCTAAAGGGCTGGGATTAC
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ACTGAGGCATAATGCTGTTAAGTAGCTTGCTCAGGGTCTAACAATTAAGAGTCAGAGCTGGGATGTGAACCCAGGTGGC
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AGAACTTAGACAAAGAGCAAGACCACAGATAAATGGATCCTTCTGTTCAGGTCTCATTACCTAGAAGAGTTTTGACTG
CAATATATGAGTACTAAAAGTTGATGGTTTATGCTAATTTTAAAGTGAATATATTTTAGAATTTTGTACATGCATTAT
TANTATGATTTTATATTATGATGGCCTTAGGAAGATTAATTTAAACAATAACAACAAGAACAACAACAACAATAACA
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GAACATGAACCTTATCTTTATAGTGACCACAGACTCCCATCTCTAGTATCATGATTTTTTAATTTGAATTAAGCATT
TTTTTGCTTTGTTAAGATGAGGCAGGCCTTCTTGCTGACATTTTAAAAAGCAACTATTTTTCTTTTCAGTTTACACTATG
AGGCATTGGCTCCAACCTGTCAGCATTGAAACTGTCAGCAGTTCCCTACCAGGAACTGGTTCCAAGGTCTAGGGTTTCC
TTAGGTAGAGGCTGGCACTGTGAAAATAATGGGGCTCTTTATCCATGTACCTGGAATGGAGTTAATACCCTGCCAGTC
TTAGTTGATTTGACATACTAACAGGATGGGTCTGAACGTTTTCTATAGTTTACTCATGAGTGACTTTCTTTGGCTTACG
TAAATGGCAAGGCCAGACAAATTAGCTTATGGACCTAGCAATCATTTCTGGCCAGATTTTGAGACACTTTTCAATCAA
CCATAGTTGCTCTAATACCTGAGATTTGCTGACAGTGCTTGGTTTCAAAAAAGGTTTCAAGTTTCTGAGCAATTTTCTT
TTATTGGGATCATCTTAATTTCTTGTGTTGCGGGGTAAAGATGAAGGAAATATGAGCAAGGACTGCACTCAGCTATTTGGG
TGACCCTTGTATACCATGAGCTTCTTAGATAGGGCCTGATGTGATCACCAGAAACATTAATTGATCGTGATGACAGGA

Fig. 9.29

GATAGTCTCTCTAGCCAGAACCTGCTAATACAAGTCCTGATGAATTGAGAATGATGCCTGAAGGAAGCAGACTGACGTG
GTGCATTAGTAATTTTAGGCCAAAAGAAGATAATGGTACTAGTATGTATATTTTTTAGTTTCCTTTTCCATTGCTTTGTT
TTGCACAGGCTGAAGAAAATAAATGTAAACAGCATATTATGGTGGCTCAGGGTGGATAATAATGGGACATCACTTCCTT
TGTTTCAGTGTGAGGTTGCCCTGCTATGTGACAGCTCCAAGGACTAAAGATTTCAATCCCCAGAAAAATGTCGAGTCC
CAGTAACATGTTTCAGGTTATCATGATTATTATGATTATAGGAGGGGAAGAGCAGTGCTGGTCTTTTAGAAAGTTCTCA
TCATGAATGTGTTCTGGAGTGAACATCACTTACTAACAGATGAGCAGCTTGAAGTTGAGTCAAACAAAACCTTTTAGTGT
TTGTAAGGGTCAGGGAGCCAGGGGACAGTCTGTAACCTCAGTTGTATATTGACACAGAGAATGTACAAAAGCTGTGAAAG
CTTCCACTTGAATGACTGCGGATGGTTGCTGGTGACGGTCTTGGACAGTAAGGGTTTTCTTCGGAGTTGTAGGAGGTGA
AGTCTTCCTATGGGAAATTTCTGGACAAAATACAAATGAAATGACTTGCAGGCCTCAGTTTAGAGTATTGTTGGCTTT
GTCTGTCAACAAATGGAGATTTGAACATGGGAGTTCAAGGGGATTTAATGAAATTTTATTAAGGAGATGAGAAGCAGG
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TTAAGTACATTAAAGTATCTAAGGGGTGTGGTGTCTGGTTATGTGTATGTCAGTTTCACATATTTTTTTTGCTTTCTTAT
TTTAGTTGCCTCTATAATCATTTTGACTTTAAATGTTTTCTGCAGATCCTTTAATAACTGCAAATGTAGAAGTATGGT
GTAACAAGTAATTGGTATGACTAACACTAAAATGTAATGGGAAATAAGGATACTATTGTAAAGAAAACAAGAAAACCT
GGGGTAGGGGAGCAGTATTGATTCTCTCTTAGGATTCCTAAGATTCTCTGTCCCAACCCTTCTACCATGGAACATTCTT
ATGTGGTCTAAGTGTCAAAGACCAGAGGAAGTGGGCAGTAACCTTATCTTCTCAATTTTCTCTCTGAACATAGATATTTT
CTTTAAGATACGAAAAACTTTTAATTGTGCTCCTGAAACCCATCCCTGTCTGTCTCAGCCCCGTTGACTCTTCTTTCT
GTGGGTGGGTGAAAAGCCTACCCATCTGCAAAGGTAGCTCTGAAACTGTTCTGGAAAATCCTGTATTTTCTCCACAAA
TGATCGTTTTAGTTTCAAGTTTATTTTCAGGTACATTAATTCTCCCCCTCCTCAGACTTCATAACAAATGATCCTGCACA
CGATTAGAATAGGAAAATGTAAATAAAATCGAAGCATATCTAGTTGCCTCAGCGACTTTATGCTTATCACTTTTCAGTC
TGCATTATTCTACTAAAAATAAAAAGAAAGATGAAAATTACCTCAGGCGTTTGCTGCCGTGCCCTTGGGTTTCTG
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TTTCCCCAGATCACCTGTCTTTTCCCTCCGACAAGGAAGCTGTGATTTTCTCTGGCCTTTAGAGGCAAAGTGATTC
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CAGGGGTACACAGCAGATAGGGCACAGTAACAGGAGAAATGTAAATGATGGCAGCAATACTTTTGTTCACTGTAATCT
GCAGCCAATTGAAGACATACTATGAATAACTAAAACATTTTTATATGAACAAAATGCTCTTCAGTGGTTCTGTTTA
TGTGGTAGAGGGCTGAATGAAAACCATGCGCTTGTTGTAAAAAAGCCTTATAAAAAGTACATTAAACACATACAGACA
CAACCATAACAGAAGAAAGTATGTGGATTGGAATTTGTGATTGGAGCAGATCAAATTAAGCCAGGGAAGCCGTTATTAG
GTTTGTATGATTGCTGGGGGGTAACTTCTGTTGCTGACAAGGTTTAGGATAAAGCTGGAGCAGATTGAAGTGGAAAACC
AGAAAACATCAGCATTTCATTACCTTCTATAGCATACTGCAGGGTAGAATTAATACTGAGTATAGACTGGTAAATGT
GAGCAGTTTACTGTTTGCTTTTAAATCATTATTGATTTCCCTTAGCCTATCATAAAAAATAATAGGGCTTTTGCTATG
AAATTAGTGCTTAGAAATATTTTTCTTCCCTCCCAATAATTTTTTATACTTTTTCTCAATACAGCACAAAGGTAGGTCAT
TAAAAATAAAGGGGTTCTTTTCAAGTTGCTTTTTGTCTAATTTTTCTCTTTAGACCTGTAGATACAAATGTATGTAT
TTGTGCTATGTATAACTCTCAAGCATAAATCATTGTAACAGTATTTAAATCACAGGCTCCTGTGGCAAATATAAATCT
TTAATAGCTATAGTTGGCAATTACTTGCCAATTCCTATAAAAAATAACATTAGTGGCTTATTTTTGATTGCACCTAAACA
ACTGGCATGATTTAGCCAGTAGGAGAGAAATATTAGTTGTGTTTTGCATAATTTTGTGTTTAGATCACACTGGAAATAC
AAAGTTTTGTGTTGTTAAAAATATTTTTGCTTTCTGAAATATTATCCCCCTTCAGATCACACCAATGAATGAAGTTGTAG
AAAGAAAGAAAAGTAGCCAACGTAGACTCCTTTTCTGTATCAATCAAATATATGCAAAATACATAGATTTTTTAAAAATG
TAATTTTAATACATTCCTTTAGAAACACATTTACTTCATGAAGAAACAGTATAATGAGTTCATTTATTGACCCAGAATA
GTGAGTTGATTTATTGAGTTTCTGTAGCCATAGACACGAATAGTAATGGTTGGCTCATTCTTAGCTATACATTCCTAAC
TGACTGTATTAGTGGGAGAAAGGAGTGCATCTATAAAATAAATAAATCCCATCCCTCAAGGTGGTCTGGAGCCCATCT
AAGAGTAAGAAGTAGTAGAGTTGAAGCTGCTTTATCTGAATCAGCTCCTGACTTCAACTCAGCTCCTCTTTTTCTTCTAG
GTCCATGTGGCCCACTTGGTGTAGTATTCACATCTCCTTCCACCTTTTTATGCCTTCATTGTTTACATTACCTTGCTTG
GGCTCTTATTGAGATGAATCACATAAAAAATGCTTAATTATCAAACATTAATCAGCTGCCTATATAATTCAGAGGATGTT
TAAACCAAGGCTCAAGGAAATCTGGTGGGTGTATACTAGGAGTATTTACGTCTTGCATCGGTACTTTCTCTGGGAC
TTCCATCACTTCAGTTCTTCTGAGATGCCAGAGTCATATCTTACTTGTGAAAGAACAGAGCTTTAAGAAATGGAGT

Fig. 9.30

CTAGAGGTGCCAGGTGCCAGTGGACAGTGGGCTGGGGGCAAGGCATGGTGGGGAAAGCAAAAAAATCTCCCAAACAG
CAATAGTTGATCTTTCTTCTATCCACTCCTCTCAATTTTCTAAAACCATTTTGTCTAACTGTGGAAGTTCTTTTGCAG
ATAAGGTTCTGTAAAACATATGCATATCTGCCATAGACAGATGGATTTGACGAAAGATGTATCCAAAAAGGAATGTATCA
CATCATGAACTTGCTAAGTGTTTCATGAATACACACTGGCATTGAGACTATTAACACTACCTACTGAAGCAAAGACAT
AAATCCAAATTGAAGAAAATATAATTTTGGATTGAATTACATTTTCCAGTGGGTTTTTTTTTTTTTTGGTGAATTTTGT
TTTTGGAAAAATTTGCTCTTCCCTCCAACGTTTTTCTGTGTATATATAGGACTGATTTGATTTCTTCAGCTTGCCCA
AGAATTTAATAATTCTTACTTTGTGTTTTCAAAAGAGTTGCTGTCTACCTTGAGCATGTTTTTAAAAAGCAGAACAAA
ACGAATGAAACAAATGTCCCTCTTCCCACAGAAAAAGCACACCACCAAGGAGTTAAATGCCCTACATTTCTTTAAGTCC
CTCCTTTTGTGGAGCTAGACACTGGTAGAAGAGGAGCTTTCATTTAATTTCAAGACAATCAGTGATTCCAACCTTAACCTA
TAACTGTGTTCTCTAATATCTGATTTCAAGAAGCAGAACATTTTGGTGAATTAATCTTAGAGTCAACGGGACCACGCT
CAGGCCATGAAACGTTTTTCTAAGCCTCAGTCTCTATGTCTTTAAATGAAAATAATGATATATGTTCTGGTATTTTACT
AGATTGGTGAATATCCACATCACAATGAGAAAGTGCTTGTCAAAGAACATTGTAATGTGTAATTTGTAATGTGCTGTA
CATGTACACTATTATTATGACTGTAGCTCATCACGCTAGGGTTAGGACTCTTTACTTCTAAAACATATTTCCAGTAATG
GACAAAACCTTTTGAAGCAAAGAAAATCTTCACTGTTTGATCCTAATGTTATGAAGGCTTTTGGACCTTACATTTGTTTA
AGCTCCCATTTGAAGCTCCCATTTGGAGCTTCAATGCTGATCCATCTATTACTTGAGTATTAAAAATACCGATAGGTTTAC
TGTGATAAACGAATGTGGCATTGTATGTGAAAACATATTTAGTAACTTCCATGTGCCATAGTCGTATAAATATTACAT
ATTGCAACAATTATCAGTATATTAATAAAAAATTTCTTTGCAAATTTTCAATTTTTTAAATGGAAAGTATCAGATATATTT
CTTATTTGATTGGATTGACTAACTTTCTAAGCTATGTTTGCTTCCCTCAAACAGGATGAATGTTCTCTGTTGGTTAAGCTT
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GAAGTTTTAGAGGACATATTTTCCCTCACAGGAAAAAGTGGGTAGGAAATTTAGACTGACAGATGTTCTTGGGTTTTTT
TTTTTTTTTTAATTCCTGTCTTGACTCTCATTATTAAATTTTGCTAAGGAAATAATTATGTAGGTTTTGAGGTGATGCT
TATTCCTAGTAGTCTTTTCATCTCAGGGATGCAATGAATGGTCAAATGACCTTCTGAATGATTTGAAAGATCAATTAG
CAGTTTTGAATCCCAAATTTTLAGGTGGTTCTGCTTCAGAAAATCATTATGCTTTTTTGAATAATGTCCATAGCTGCAT
CCTACATTTCAGTGGTTTCAGCTCACTGCTGTTATAGTTTGATGATTTCTTAACAGCAATATTGGTTATGCTAAAGCAG
TTCCCATTTTCACTGTTGAACCTTTTTTAAAGATAGAAGAAATTATAGAGGAATCACAATAAGTAAAACGTATTAAA
ATGGAGTGGATTATCTCCACTTTTATCTCACACAGCTCACCCAGAAATTCATGAGAGAACTTTCTAGGAATGAAACAAT
TTCATTTGTAGTAGTATTTGAAAACCTGGATCTAGGGCCATTGACCTGACTTTTTTGTGCCTTTGGTGATTGGATAAGAA
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AATCAGGAGTTAGGTGCTAGAAACACAAAGATGAACAATACATGGTTTCTCAAGGAGCTTATAACCTACTAGACATTT
ATTTTCATGTTGGCAGAACTTTTAGGATAATCTCAAAGAAAAAGGGTTATTATGAGGTGCTACTTTTCTCCTCAAAT
ATTTTCTATGTTAATAATGAAGGAATGACCAATCTGTAGTATATGCAAAAAGTACTGGGTAGAAATATATTAATTTCT
TGGCTGGGTGTGGTGGCTCACATCTGTGGTCCCAGCTACTTGGGATGCTGAGGTGGGAGGATCACTTAAGCCTGTGAGT
TCCATGCTGCAGTTGAGTCATGATTATATCACTGCATTCCAGCCTGGGCAGGCAACAGAGTGAAACTCTGTCCCCCACC
CACCCCCAGAAAAGAAACATAAAAAATTTTCTTATAAAATTTTACTTTTTTAGAGTCTCCAGCCTTATTTAATTCAGTTAC
CATATTTGACAAATAAATGGTGGAAACCAAGATGTTACCAGGTTTAAGTTTCATATACAGTAGAACTGTGACAGGAACCT
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GCTAAAACAAAATTTATAAGCATCTATATATTTTGCAGCTAGGACAATTATCTACAAACATGATATTTAATGGAAGATA
TGGTAATAACATCTGAGATATACAATTATGTTATAAAATCTAAAATCAACAAGAAAGGAATAAAGTTGTACCATTCCAG
GAAATACATTTCCAGAGCTTTAGATATCCTTATTAGATTCTACATGTTAGTTTTGGTGATGTTAACTGCCATAACATAT
AAAATCTGAAATCTTGTGACTTAACTGAGAAGTTTATTTCTCACTCAATCAAAGCCTAAAGTGAGTATTCTTGATGGG
CAGATTCTTCTCATTCAATCAAGCAGTGTTCTTGTGACTCCACTCTTANCACCTTGTTGTACNACCAGCTTCCTGGTG
TCTGTGCTTTGGCAAAGGAGAAAACCTCATAGAAGTCTGCTCACCTTCTATTGGCCAGAGCTTGGTCACATGGCCCCATA
TAGTGAAAGGGAAGCTGGGGCTATAGTCCTATTTGTGTCTGGGAAGAATAAAAAGTTGGTTTGGTGATTGTCTTAGTTC
AGGCTGCTGTAACAAAGTATCATCAACTGGGTGTCTTATAAACAACAGAAATTTATTTTTTTTTTACAGTTCTGGAGGC
TAGAATTCTGAGATTAGTGTGCCAGCCTGGTCAGGTTCTGGTGAGGGCTGTCTTTTGGGTTGCAAGTGGCTGACTTCTC
CTATTATCTTCACATGGCAGAAAGAGGGTGAGCTAGCTCTCTGGCTTCTTTTTGTAAAGGCACGAACCCCATTCCTGAT
GGCTCCACTCTCATGACCTAATTATCTTCCAAAGANACCATCTCCAAGCACCACACATTGGGAATTAGATCTCAGCAT
ATGAATTTTGGGACGATATGAACATTCAGTCNACAACAATGATCCACTAAGCTATTTCTGCCATACCTTTTTTTCTTTT
CTCTTTTCTTTCCTTACTGTTCTTCAGTATGTCTATGTAGAATCTATGCCTTGTAACCAATAAGGTAATGAAAAAATT
ATAATATTTACCATCTGGAAGAAATTAGGAAAATGATTTTACTTTTAAATTCCTTGATCCATATGTTAAAAAATTTCCAA
GGCTATTAAAAATATCACTGAACATTGTGATTATAGAGATTGCTATGATTTAATACTTATGTCTTATAGTGAAAAAGTAG
TCCTGGGATATTTTTCTCTCTTTACCTCACCCATCCTCTTCTATGTCAGGTATTTGAATCCCTTGCTTTGTAGTTTTAT
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TTGGAGAAATTTTTCTTGCATGGAGTTGAGCAAATGGTGCACTAACATAAGCTTGGTTTATAGTTTCTGTTTTTTCAGA
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ACCTCTCAGTAAACAATTTTTTTTTTCAGATTATTCTATTTAAAAAATCTCTCCAAATCTGTTAATTTTCTGTTAATAAAA
CATCAGTTTCAATAAAAGTCATAAGCCTGTCAAACCAATTTTAGTCCTTTATCTTGGATATAGTAGGTTAAAAATATTTT
TGTGGGTGTTAAGCCAATAATAAACTCAGCTATTTTAAACAACCAATTTATTTTCTCCTATAAACCATAATTTACAATC
AAGCTACTTAAGAAATAAAGACAACAAATAGAACTGTTTTAATCTTCATATCCTGACCTAAAAAATAGAAGCACGAAC
TCATGTCACATTCTCATATTTATCTGATTAGACTAGGAATGCCTAAAAAGAATTTTCCCATAACTCCATTTACTTGTG

Fig. 9.31

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TTTATTTCCCTAGCTCATTCCAAAAGCAATTTAAAATACAACACCAAAAGAGTAAAATATCCTCAATCTGACATCCTAG
CAATCTTTTTTATTTTGGTGAAACTGAACAATAATAATAGGCACTTGCAATTCCTCACCTTGAGGCACTGAAAACACTGAT
ATGTCTCAGTTTTAAGATTCTGAAGATTATATTCTATTTAATAGATATTTTAAGTATATGTATTATATTTCTAGTCTTT
GAACTAAGGCAGTCCACATATAGGTATAAAGTAATGCCTCAGCATAGTGTAGCAATTATTGGTGCAAAAGGTTGGTGCA
AAAGTAATTGTGGTTTTTGTTCATGAAAATGGCAAGAACTGCAATTACTTTTGCATCAACCCATAACTCTGTGGTCTCTG
GAGCCAGACTGCCTGGGTTTCTAGATCCAGCTCTGCCACTTAGTACCTGAGTTACTCTGGACAACTAATTAATCTTCAT
GTACCTCCATTTAATTATCTGGAAAATGGGATAACAATAGTACCTATCTGGTTGGGTTGTTGTAGAGATTAAGTAAATT
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AGTCAGTTTCAAACCTTCTAGATATAGCCAGTTTTCATGGGTCTTTACAAAAGGATCCCTGTATCTTTGTTTCTTCTGT
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TTTAGGAAGAGTGTTTGCCCTGCACTAACTGGCCAGACTGCTCTACTGCATCAGCCCTCTGGCGATGTTCTAAGGGT
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CTTTCACACACTGTAAGGGCAGAATTGAGTGGTTCCTCGCAGAGACCATATGGTCTGTATAGCCTAAAGTATTTAATATC
TGACCTTTTACAAAAGAGTTTACCAAACCTCTGTTCTAGAGCTTTAAAATTGGACAGAGACAGAAATAAATGAGATGTTA
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AGAATTTATTGCAGTAGGCTATGTTGGATAAAGAACAACAGCCCTCAGTAGCACATTGTGAATATTAAGAGGTGGTGA
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ATGCACTTTTCAAGTCTCTTTGTGTGTACATTTGCTAACACCCCACTGGCCAAAGCAGTTTGTGGTGGAACCTCCGAGT
CCGAGTGGAGGATTGTAAAGTTATGAGTCAAAGTGTGCATACATAGGAAGGCCATGAGTTGGGCCATTAATACACTCAG
TCTACCAGAAGTCATTTACCTTTTTTCTCCTCATCTTAACAGCCTCCCTTTTGGAAATTTAACTTTTACATACTGTGTA
AAATCAAACCAAACAAACAGAAAAATCTCAACTTACCTGGTTACTTTTTAGCACTGTTTTTAAAGCTTGGTCAGGGAGT
AAGTTGTTATTACAGGCTAAAATTTAAATTTTGTCTTATTACTTACACACTTCTCCCTAACTCCTCTTTTAAAGCA
AATTGAGTTTAAAGCCACAACTGTTGAATAATGAAGTTGGTTCTTTTTTCAAGTTTATTATCATATCTATCCCTGAGCTCTA
TTCTGCCTCTAATCAATTTCTGTATTTTGGAGCCGCTATTACTATAGCATTGTGCTGGGTGAGTCCGTGGGGTTTTGATC
ATGCTTTTCATATGTTCTTGCTGTTTGATTTCTCTCTCCTTAACTCCAGGTTTGCATTTCTTCTGCTTATTCTACATG
TCATAGAGGAACCTCAAACCTGGACGTATTTTAAAGAATACTGGTTATCCACTTTGCCCAGTCACACATACCTTCAACCA
ACAAATATATATTTGTTGAGTGACTTCCATACTATTTTCAAGGCATTAGTAATATACTAGTGTAAAAGTAGACAAAATGGA
GGCCATCCTAGAGTTTATAATCTAATGGGGATTGACAGACATTAAACAAATGTGTGAATACATAATTTAATGCTTGGCC
GTGATAAGGGTTATTGAGAAAATAAAGCAAGGTGAGCAGAAAGGGAATGATGGCGGAGGTGCTTTTTTTTGTATTAGTGTT
GAGGGAAGGCTTCTGAAGTGAGGGAGTGAGCCATGACAATCTGGGCAAAGTGTGCTGCACGCAGGGGCATTAGTGGATG
CAAGGCTGTCTGAATGACGTTTGGTGCTGCAGCAGCTGTACTTCCAATGGAAAGAAGGGTAGGAAATGAGGTGGGGGAG
ACAACAAGGGACCAGATCACGCAAGGCCTTTGTAGACAACAGTAAGCTCGTTACATTCCTTCTAAGTGCCATGGCAATC
CATTGGTGGATTGTTGGGGCAGGGGAGAGACATGGTCTGATTCTGGCTGCTGCGTGGGGCACAGGCTATAGGGAGCAAGGG
TGGAAGTTGGGAGGCTATGACAGTCATGGCAGGGGCTGTGTTAGGAGAGTAACAGTGGAGATTGTGAGAGGATGGCAGA
CTGTGGAGCTCTCAAATGTGTTTCAATTTAAAGAAGCCAAATTTAGATACTGTGATCAGTTCTGCCAAGTCGAGCCTG
TGGAGAAGTATCTCCCAAAGTAGACCCAAACAGACTTCCGGTGACATCTCAGTTACTTTTAGGTGCTATGTACATGCAT
GAACTTTTAAATTTAATAGTTACTGATGATTTTAAATGTGGATTAGAAATATATAAACTTGCATGTGAAAACCTATGAT

Fig. 9.32

Fig. 9.33

Fig. 9.34

Fig. 9.34

Fig. 9.35

GGTGAATGTTTTTTTGTGCTGGTATGTTCCACCATGTGTGTGAGCATGAAGGTAGGAAAACCTGTTTGCTCAGCCATT
CCCTCTGTGGTTTGGTGTGTTGGAGAATCAGTCAAATGGAAAGGCTTTCTCTGATGTGATTACAGAATCAAATATTTT
ATAATTCCCAGATACTTCTGAATCTTGAATAATTATTAAAGTCAGTGAACACTTTCTCTTTCTTTTCTTCTGACTCCCTT
TGGATTGAACTAGCCCGCAGGGCAACACAATATTATTTCCCTGGGCTAGGTGCACCTTTCTCTTCGTGGACTATTGGGT
GCCTGTAGATAGAGTCCCTGTGCACAGCAGGCAGATTGTGCACCGCATAGTCTTGAGGTAATTTACCAAGAGTATA
AGTGGGATTGTGCAAAATGGTAGCCCTGCCTATGCATGTTTGGGGCAGGGATTCCCCCAACTGTGTATTTCTGGGAG
GCCTGATTTTCTCTCTCTTTCTGTCAACCCATCATAAGCATATGGAGAAGATGGATGTTTGAAGGGAAAATAATGTTATT
TTCTGCATTTTCTCCCATTTTTTTGAGAGCACTCATTCTCCTTCTAATATTAATTTAGATCAGGGAAAATAAAAGCCATT
TGTTAAAGAATACTAATTATATAATAAATTATCAAACAATTACGCCAAAGGTATGTTTACAACTCCTGTTCTATCTT
GGTGGGTTGATTAGCTTCTTGGGGGGTCCAAACAGCTATGGCTTTTCCATTTTTATAAATGGTTCTGCATTTATAAAA
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GCTCAATTTTAACCAATAAATCTTATACCAGAGTGTTCTGCCTGCTTCCGGATGAGCAAGGAGACTTGTGGTCTGGTTC
TTGGACTTCCCTAGCAGCATGGCCCCCAAATGCCAGTCTCCACTCCTGCCTCAAAGAAGAAACCAAGAATGCCTCCTG
CTCTGGGACCGGAGGAGACATTGGCCTCTGCAGGCTTGCTGAGGAAGGGAGAAAAGAACAGCAAGAAGCAATTGAATG
CATTGATGAAGTACAAAATGAAATAGACAGACTTAATGAAGAAGCCAGTGAGGAGATTTTGAAAGTAGAACAGAAATAT
GACAACTCTGCCAACCATTTTTTTCAGAAGATGTCAGAATTGATCGCCAAAATCCCAAATTTTGGGGTAACAACATTTG
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ATTTGAAGATATTAAATCAGGTTACAGAATAGATTTTTATTTTGATGAAAATCTTTACTTTGAAAATAAACTTCTATCC
AAAGAATTTTCTGATGAGAGTGGTGATCCATCTTCAAAGTCCACTGAAATGAAATGGAAATCTGGAAAGGATTTGA
TGAAATGTTTCTGGAAGGATTTGATGAAATGTTCAAGTCAAATGCAGAATAAAGCCAGCAGGAAGAGGCAGCATGA
GGAACCAGAGAGCTTCTTTTACCTGGTTTACTGACCATTTCTGATGCAGGTGCTGATGAGTTATGAGAGGTCACCAAGAT
GATATTTGGTCAAACCTCATTACAGTACTACTTGGTTCCCGATATGGATGATGAAGAAGCAGAATGAGAAGAAGATGATG
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TGAAGGGGAGGAAGGAGAGAAGGATGAAGGAAAAGATTACTAGAACACTGATAGGTTCCAACCTTCTGTTTAAAAATT
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CACTGTAAAGGAAAGATGATGCTCAGTTTTTAAACGTGAAAAGTACAAGTTGCTTTGTACAATAAACTAAATGTATAC
ACATACACACACACACACACACACACACACACACACAACTTATACCAAATCACACCCTTTAGTTAGTTGTCCA
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GGTTCAAGTGATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGACTACAAGCATGTGACACCATGCCAGCTAATTTTTT
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TTGGCCTCCCAAAGTGCTGGGATTACAGGTGTAAGCCACTGCACCTGGCCATGACACATATTCTTAGCTATTCTCCAAA
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GATTCTGTCTCTCAGCTGTACTGCAATCTATTAGGACAAGTCTCGTCTCTTTTATGTATTCTTCTACTGTGTCTATA
TGTGGTGGTCTCTCAGTCGAGCAGGCTGTACAGTTTATTTCGAAGTGTTCCAAGGAGCTTGTGTGCCTCCACTGCTCTGA
CCTCATTTTTTGCAATGTTAATTAAGTTGGGTAGATGACCTATTTCGACTGATAACACTCACTAGCTCACTAGTATATGGT
AAGTAGTTTATTCTCTTAGTTAGTTGACCGTGACCAGATTAAAGCCCAAAAAGACAAGAAAAAAAATCAGGTGGTAGAG

Fig. 9.36

AGTGGGGAACAGAGGTATAGGACACTGTTTTGGGAAAGTGGATCTTCTTGTTCTTACAGAGGATAACTCGGGCNTGT
CAGGATGAGGTATTACAAGAAGAGAGAACAAGAGNGGAGGCTCACCATTAAATTTGGCTTCTGGCCTTGGGTCAGCCTT
GCATGATGTTTTGTAGACTGAGAGTAAGGAAGGAGAAGTTGAGACAAAACTTTACTCTCTGTTTGGTCACTGAGAGTA
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CCCAAGATGACTTCCAGGGCCTACGTGGGTCTCTCTTGAGTTCTCATCTCCCGAAGGTGAACCTGTACCACTTGTGTTGA
GCACTCCTAGGACCAAAGAGTGTGGGGCTGTGCATGAATCTTAGATGTTTGGGCTGAGATGTCCTTGGGGGTGTGTGCA
AGGCCCCCTCACTGCAGAAAAGAACTGAAGGTGGGGAAGAGAAGAGAAGGGAGTGGGCTACTGGCTGGAGAACCCTGGCT
CCAAAAATTTTAGGCATGGGCTGACATTTCTAACTCTCCAGTGGGAAACCATTTAAAGCAGTCAACACATTGACTACGA
AGTGCAGGCACTACTTTGAGTAAGTAGACTTGCATGCAGAGGTCCTTGAATGGCCTCCTGGTCTGCATGTGTAACAACA
GTGGTTATTGTTCTCTGGACTCTAATGCTGTAGGTACTAGATGTGACTGTTCTCCAGAGACATTCCTAGGGAGTTGCTA
ATGGTGGTGGTGGTAATTGCTATTATTATTTTCATGAACATTGAAATCTTTTATTTTTTTGTAAAGTTGTCTCCTTGAAT
ACTGGACTGTAAAGGTGTGTGGATGTGTTCTAGCAAATTGAGAATTGTAGTTAATTGAGTTCTGGTTATTTAACTTTTT
ACTGTAGTTTGTACTTGTGCTTGTTCAGAGTCTAGGCAATTTTACATGGGCTTATTTTATCCTTTCTCCCTATCTCCA
TTAGTCTTCAAGATCAGGAAAATGATTTATGTCATTTTAAATAGAGCCTTCTCTGTCTCTCATTTTCTCTCTCTCT
CTCTCTAAGGTATATTATTGCCTACTATATCCTGTGCCCTGTGCTGGAATTTCTCTTGTGCTAATTTTTTATTGATAACA
TATGATGGCTTAAATGTCATCTAGTGTTTAGTAAATTCATAATAGAAAATAACTTTGTTCCTCACAAATTTTTTACAG
ACATTTAAATATTAGCACTTCAATATCCAAGTATTTTGCTTTTTCTTTTCTGGAGAGAACTAACACCTGTAGGTATTTG
TTGACATGCAAAAAATGCATGCAAGAATTTTAAATTTTATAACCTATAAATTGGTTGTATACTATTAATTAGACCTCTT
CTTTAGCTCTCTTACTGTATACACTCTTCTGTTTTCNCAAATCTCTCCTCTTTCTCTAAATTCCTCACCCCTGTTCCCC
ACCCTTTCCCTGCCTTGTCTCTGGGTTTTTGTCTCTCTTGCCTCATTTTTCTCTCTCTCTCTCACTTTCTTCCCCCA
CCCCTTCTCTCTTGTTTACACTCCCCGCTCATGTATGTGCCGCTTCGTCTTTGTGTTTTATATCTACCAGGGGAAAGACC
AACCATCTTTATTTTTTAAATTTTATTTTACTTTAAGTTCTGGGATACATGTGCAGAATGTGCAGGTTTGTTACATAGGT
AAATGTGTGCCATGGTGGTTTGGTGCACCTATCAGCCCATTAACCTAGGTATTAAGCCCCGCATGNGTTAGCTATTTATC
CTGATGCTCTCCTTCCCTCACTGCCTCCCAACATGCCCTGGTGTGTATTGTTCCCTTCCCTGTGTGCATGTGTCTCAT
TGTTTAGCTCCCACCTTACGAGTGAAAACATGTGGTGTTTGGTTATCTGTTCTGTGTTAGTTTGCTGAGGATGATGGCT
TCTGGCTTCATCCATGTCCCTGCAAAGGACATTATCTCATTCCTTTTTATGGCTGCATAGTATTCCATGGTATATATGT
ACCACATTTTCTTTATTTCAGTCTATCATTGATCGGCATTTGGGTTGATTCCATGTCTTTGCTATTGFGAATAGTGCTGC
AATAACATACGCATGCGTGTATCTTTATAATAGAATGATTTATAATCCTTTGGGTATATACCCAGTAATGGGATTGCT
GGGTCAAATGGTATTTTCAAGTTCTAGATCCTTGAGGAATCACCACACTGTCTTCCACAATGGAACCAACCATCTTTTAA
AGTAACAAATGACATCTAAGTGTGAAGTCCGAAGTCAAAGAGCTAGAGAGTCATACAGTTTCAGAGTTGTGCGAGTTTT
GATGATTGATTCTCTGGCAGGGTCGTCTCATTTTGTACCTGAAAATATAGTTCTAGTGAATTATATCACTTGTCCAAG
CCCACACAATGACCAGGCTTCTAGCCCTGATTTATCAGTCCAGTCCTTTTCTTAAAAATAGTTTCTGAATATATGCAG
GTAAAAATAATAAATAAGCCATCATAAGTTACAGACATGTATGTCTGCTTCTATGGTGCCAAGGAAAGAAAACAAGGGG
AAGGCATGAACATAAAATAACACAGGTGCACCTTATACATTGTTTTCTTCAAAGATCATGAAAGGATAGCTTAGAAATTG
CCAGTAAGAACTATGACAAACAAGCTCAGACAAGTTACTGCAGAGAGGAGTGTACTTAAAGTTAGAAAGCGGGAGAAGT
ACACTTGTCAACAGGCAGAAAAGGAGGAGCTATGGACCTTACAATTGGAAAATGTGATTAAAAAATAACGAAGCT
GATCCTTCTTGATTTTTCTTGTTTTGAAATTTATGGCAACCAGTACAACAAACCTATTTGAATGTATAAATTAAATTA
GTATACTGTGATTTACTTGGGCTAATAATAATGTAAAACCTTTAAGCTAGACCAGCTGTAATTATATTTCCCAAAGATT
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GCTTTCCTGACTTGAAGTTCTCATGTGTGTTGAAATCGAATTAGGAAAATGAGTAAGTAATAACTTATCCTAAGAGTA
TAATCTCTCCTTTGTGACATAAATTAGAAATCCATTCACATCTGTGGAATAATTCCCAGGTTTGGGACTACACAATCTG
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TGTGCTTTTAAATAGATGTATCATTCTCAAATTTGGATTATAAAGAGCTGATCTGAGCTGCAATCAGATGGATAATT
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AAAGGTAAATTTGACCAATGCACATGTACAATGATGTAAGGTTCTCTCACTTTCCATTTAATCTCAGGATTGTGGAGGC
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GCTTTATTGTCTCAGATCTGTTTAAATCACCTGGAATACTTTTTCGAGTGATTTTTTAAATGGCAAGAAATGTTCCCTTT
TCCATAATGTAGCTGTTGCCCTCTGAAATGCCATAGGTGTTTGCATTTTTTAAATCCTCATTTGTGGCAGTTATTACGTT
TTGCTTCATGTTATGGTAATTTGGCATGCAACATATCTTTTGTACCAAATTTTAACTCTTTGAGGACAGGGTTCATA
TCATTGTCTTCTTTACCCTGGGCTTTTCACTGGGTGGTGAAATGTTTGTGGAATCGGTGAATGAAGATAAGACATA
TTGTCTTCTGAGAGTGGGCAGACATGTGGCGGGCTATTGCTGGATAATGTGCTCATAATATCCTGCAGATTAATTTGGT
TCTTTGATGGAAAATGAAGTGAACATGCATCTGCAGTCCAACCTTTTCCCAATGTTATTCAAATGTAATTCAATGT
GATCAATTACATTGAAAATAATTGAGAAATGAGATGACACTTAAATTAGTTGAAGACAATTGTGGATTTTGCCTTGGT
TCTTTGGATAAGTAACTTTTAGTGTTTTCTCAGAGATTTACAAAATTCAGGGGCTGTGTATTCTTGCCTATAGGACA
TCCAAATCCTGAGGCTTTTCTTCTTATCATTTCTCACTTAACTGCTTATTAATCTTTCCAAAGAGAAGGAATAAGAAAC

Fig. 9.38

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CATATGATGATAGTTGTAGATTTTCAGAGGCCCATATGGACCATTTGAAATGAAAATGTTATATTGGCAATGGATGAG
AACATGCCACTTGCCCTGCTCCATACAACTTTGATTCCTAGCAATCAGCACATGTTAGAGAGCTGGAGGCCACGGTAT
GGTGTGCAAGTGCTCCTGTGCTGAAGTGACACCAGAACAATCACCACATTGCCATCAGGCCGTCCATCTGAGGTAGGGT
TTTCCTTCTTGTTCATCTTTATCATTCCTGGAGAATTTGCCAAGGCCCCCAGGGCAGTGCTGAGGCCTCCAGCGCCC
CCTACAAATCTCTTTTTCTGTCCCTTCAACATTTTCAGGTCTAACTACCTGTAGTTTTCTCTCTAATTTCTGGGCCAA
AGCTTGGCAGGGGGCAGGCGCAGAATTAGGTTGAAGAGGCCAGTATGAGCAGGCATACCAAGGTCCATTGTGCTATACC
AGAGGTGTTACATTAGACTAATGAAGGGATTATAGAACTTAGCTCTCCATCTGAATTCTCTTATTAACACATCACTGT
GAATCAGCCTGCTGCTTTCGGTGTGTGTATTTTTTCCCTTTGTTACTTCTGTACTGCTAACGATATTGGTATCATTT
GTTGTATCAATGCAACTGACTTGTACTTCAAATTAATATCTCAACCAACCTAGAATTTCTTGTCTAGCTATGGTAGGTA
GTATGGGGTGGTAGTGGTGTGTCTCAGGCTAGTATATGATAATATGATTTTGAAGTCTGTTTTCCCCCCCATTAAACG
AACTAGTGTGAACATCTATCCATGCCTTTTAAACACTTTTCTCCAAGATAATTTTTTATAGTTTTTTAATGGTTAAATA
ACATTAACCTTAGCCAATGAAATTTAAATATTTTTGCTACTGTAAAAATTATTGTGTATCTCCTGGTAGTTAAATCAGGA
TTATTTCTTAGGATGCAGAGCTTGTATGTTAAGGAGTATAGTTATTCATAAAGGTTTGATTTTTTTGGCCAAATTGCT
CTCCAGAAAGTTTTTTTCCCTTCAGTTTATTCTCCTACCAGCAGTGCATGAGAGGGCACATTTTCCCGTATCTTAAGT
AGTGTGTGTTTTCTAAAAATGTTGCCAATCTGATATGGGGAAAATGGCATTTCAGTGTGGTTTTCTGTGTTTGTAAAGT
CACTAGTGAGGTTGGGTAGTTTTATCCTTTGCAAATTGCCTATCAATGTTCCCTTACTTATTTTTCTTTTGAGATGCTTA
TTTTCCAATTAATTTTACTGACAATTTTTATATAGTACAGATAGTAATCCATGTGTCTATAGGTTGTTTATAATGTAAA
TATTTTTTTCCAAGTTTCTTAGTCACTTTTTGACTTTTTCCCTTGGGTATATTTTTACATGAGAAGTTTTTGTGCATGT
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CTGCCTTTGGAGTTAGTTTAGAAATGACTTCATCATCCCAAGATCACACACACTTTTCCCTACATTTTCTTCCAGTTTCT
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TCATCATCTCCTGACCAGATGGTCGTCATAATGTCTTAATTGCTCATTCTACCATTGGTCTCACTTGCTTCTAATCCAT
TCACAACAAAGTTGGCTGAATTGTTGTTCTAAAACACAAATTGGATCACGTTACGCCCTGCTTGTGTGTATACATACT
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TGGCAGGACATAGCCCTGGTGAAATCACTGGCAGGGGCTTGTGTGAAGTAACTGCATAAGAAAAAACAGCATGCTGTT
AGAGGTGAAGGGACCAGCTTTGAGTTTAAACAGAAGTAAGGACAGAATGGGTGAAGCCATAATGTCAACTTTGAAATATC
ACCATTTATTAAGAAACCACATTTGAAGTTCTAGGATCCTTAAATGAAGGCGCTTTTCACCAAATACTTTTCTAAATTT
CCTGGGGTGTTTATAACATGGAGGACCTATTTCTGAGCTACCAGAGTCCTAACACATACCTAATTATGAGTTGGTCAC
CAAGAGTTATCAGTAATCACAAGTAATGAAAGCATGGCTTTGGGTGAGTGGGTGGGTAGGGTGGGTGAGGTGGGATATG
CCATGTTTTGGACCAAAGCTGTCCACATTGATGTGTAGCCATTGTTTTCATGATTCTAGCACCCAGTGGATGTTGAGTA
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CTCTCTGAGATGGCAATGCTAATTTTGCTGGAGAAAGTAGCATCACTGTTTTACTTGGTTCCCTCTCTTCATACTGCCT
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CTTTCAAATAAATGTTTAATTTGAAATATTTACATTAATGATGCAACCATTTTCTATCCACCCAACTTGAGACCTAA
GAATGAGTTGTTTGTGCTGATTCTTCTCTCATAGCCAACCATTTCAATGTTCTCTGATTTCTACTGTTACCCAGGCTTTGC
TTTACATTTCCACTTTTATGGTGCTTATCTAAACCCTTACTTTTGGCCTTTAATATTGTATTGGCTATTTATTTATTAT
TTTATTATTATACCTTTAATTTCTGGGATACATGTGCAGAACGTGCAGGTTTGTACATAGGTATACACGTGTTGTGGTG
GTTTGCTGCACCTATCAACCGTCATCTACATTAGGTATTTCTCTATCCCTCCCTAGTCCCCCACCCTGACAGGC

Fig. 9.39

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CCTGGTGTGTGATGGTCCCCTCCCTGTGTCCATGTGTTCTCATCGTTCAACTCCCACCTTATGAGTGAGAACACGTGGTG
TTTGGTTTTCTGTTCTTATGTTAGTTTGTGAGAATGATGGTGTCCAGCTTCATCCGTGTCCCTGAAAAGGACATGAAC
TCATCCTTTTTTATGACCGCATAGTATTCCATGGTATATATGTGCCACATTTTCTTTATCCAGTCTATCATTGATGGGC
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TGATTTATAGTCCTTTGGGTATATACCCAGTAATAGGAATGCTGGGTCAAATGGTGTATCTGGTTCTAGATCCTTGAGG
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ATCCACTCCAGCATCTGCTGTTTCCTGACTTTTTTAATGATTGCTATTCTAATTGGCATGAGATGGTATCTCATTGTGGT
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TGAGAAGTGTCTGTTTCATATTCTTCACCTCTTTTTTGATGGGGTTGTTTGGTTTCTTTCTTGTAATTTGTTTAAAGTTC
CTTGATAGATTCTGAATATTAGTCCTTTGTGATAGATTGCAAAAATTTTCTCCACTCTGTAGGGTGCCTGTTT
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GCTTTTTATGGTTTLAGGTCTTACGTGTAGGTCTTCAATCCATTTTGAATTAATTTTGTATAAGGGGTAAGGAAGGGGT
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GTGCTTTAGTTGCCCTCGGTGAAAAATGGGGATAATCATAGTGCTGCCTTATAGGGTTGTTGTAATAATTAAATGATTAT
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ATCCAACCTTCTCTTTACTCACTATTTCTAAACACATTGTGCATTTTTCAATATATATGCATTTCAATATCTATCTTTT
GCTTCCTCTGCCTGTAATACCCTTTTCTACGCCCTATCTCTCCTGTGTCTTTAGAAATTCTATCAGTTCTTCAAATGAA
CCTCAAATAGACTTCCTCCACAATCCTTCACCTATCATACCAGGATTGTTTATGTTTCTGAAATTTACAGCCTATCTT
ATACTTTTCTGTTTACTTGATTCATGTTGGCTTAGTTTAAATTACACACGAAGAAATCTTGTGTTGCCTGATGGACTGT
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TGGGATGTCTATTATCCTGTAAAAACATGTATTTCCGCAGTTATAAGAGCTTTGATATTGGTGTGAAAACAATGGTCC
CTAATTAAGCCATAAGGTATTTTGGGAATTTAAACAATGTTTGGTGGTTATCAACTGGGAACCTTCTGGAGAGGTGAAA
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CATTCTTCAACCTCTACAATTATGAGAAAGTAATGACTGTCAAAAAAAATTTGCTCATTTTTTCTGACATCTTTCAAG
GCCCTGAATAGACGTCTTAATTTCCAGGTCTGTGTAGACTGACTTGGGTACTCAGGTAATGGCCTCTCTTCTGTTGAT
GAACACACGTATTGAAGCATATATATGCCAGGCTCAGTACTAGACACAAGGGGTACAGTCAATGAACAAAATACACACA
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GGTTAACAGGAAAGGCATTTCCAAGTACTTCTCCCTGCTTCTTTCTCATCTGTGTGCTCTAGTTTTACTCTCCCTCCCT
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TTTAAATCTGTAAATTTATAGGGCATATTCCAGCCCTACCAAGGGTCACTTCACTCCATTCTCAGATAGTGCCAC
TGTCCTGCATTTGCCTGCTGTCAATCACAAAAGAGATGGCTTCTTAATGAGGCTGGTTTATGACTTTTTTCTTAAAGAC
AAAGTCAGTCTTCTTTTGCAAAAGTTGTTCTAATGAAGGAAAGTAGAAATAAAAATTCATCATAGCTTCTATGGCCAGA
CATAAAATGTCTACAGCAGGATTGATATGGAAACATAAGCTTTTATTAATTTGTTTATCTGGATGCTTCTGAAGCTAGA
AAGAAAGACTTTCACCATGAAGCACTGGCAACTGAATTTCCATAATGGATTCAAAGATCTAAAGAATATCCTCAATTT
ACATATCTTTCTGAATTTATGTTCAAAGAAAGGGTATTAATATAAACATAAAGTAAATAAAGACAAAGCCCAAGTGTA

Fig. 9.40

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TTAAGACTATTCAAGGCCTGCCATGCAAATTGCAGTGGAAAAGTCTACCAGTAGGTTATTGGGAAACAGAGAGAACTT
GGAGTCCAAAGATAGGAGAGGATACCCACCCTAACTTTCAATGAAACCACAAGCTGGCCACTTAGCTTTTCTTGTGCT
GTAGAGTCAAATGTGAAACAATAGGCTTTATTTATTTAATACTTTGTATAAGGCACCTTTGCTCAGTACATTATATATTG
TCTCATCCTCATAGTGAGCCTGAAGGGTAGTAATTTCAATTGTACCTTACAGATGAATAAACTTAGACTTAGAGACCGGT
CAAATAGCCTGCCCAAACAACCAGTTAATAAATGGAATTGGGATTTAACTTAGATTTGTCTGATTCCAAAGCTTGTG
CCCAACCAGTGTTTATTGAGTACCTTCCAAAGAGCCAAGCAACAGTTTGAAATAGGTTTCACATATGTTCAACTATCACA
ACAACCAGATAGAGTAGATATTCTGTTGAAGATGAGGAAATGATGTGGGGGAGAAAAAGTGATTGTGTGGGGGAGAAAA
AGTGATTGTGTTGAGGTCACACATCTAGAGTGGTCGAGTCAGAATATGTGTGAACTCAAGACCTCTAGCTCCAAACCTA
TGAATTTTGTGTTTTTAAACATGACTATAGTCTATTCTCTGTACAATGTTTCATAATATTTTTTCTGCCAGTTGCCTGCTC
CCTTTAGGTAAAGGGATAATTAGCAGAGATTCCTTGCAGAAAAGTGTCAAATATCAGCACTTAAAAACAGTGCTGAATT
TTCCCGTTATTTAGCCCATTACTTCATTGAGCCTAATTTTCATGCTTCTAAAAACAGCATAAGTTCTGTGGGTAGTGTTT
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GCCACCTGCAAGTAGCTTAGGCAGTGGGCAATCATACTTCAATTCTAAAGGGTGTGGAAGGGATGAGTATTTTCTGTTT
GAAGCTCAAGCTAGATTAAATGAATTCTTGATCTTAATCTACTTTGAACTCTACTTGGAACACATCATGAGTTGTTTTG
GCTTACTATTAATTTTTTAAATACAAATACAGTGCTCTTGTATTGAAAGCATAGGATTTTGTGTAGGAGAAAAATTTGAAT
GTCTTGAAACATTCAAAGGAGAGCTCAGGGAAATAAATGAAAGCTGGTGGATTTCTAAAAATCTTTATTGAGGATTAA
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CATTTCCCAAAGATTTTTTAAACAGCAGAACCAGAAAACTAATTCACAAATTATCAATATTCAGCACTTTGATAAATTC
AGATGAATTTGCCTACCTAAAGAAATGCTACCTGAAAAGCTTCTCTGGCAGGCTTCTCCAGAGTTTCATTATCTACTCT
GAACTTTTGTGTTTACATAGCAATCATTTCTGCTGTGTTCTTTTCCCTACTAGCCTTGGTAGGCTTCTTAGCTGAATTGT
CCTGAAACCTATAAACCAGCCCTGGAGGCTCTGAAAAATTAAGATCTGTCTTTTTTGTGTTCTGTTTAAATAAATGTTAGC
TTAAGAAAGTCTGCAATGGAAGAACACTCAACTTGTCTAATGATTCTCTCATTATTCAAGTTGCAAATAGAAAGAG
GTATTTCCCAATTTGCAATAAAATATTGGACTTTAATTTCTAAAAAAAATCTTGAAAGCTACATACTCCATAGACTTC
TAAGCAAAGAGAGTCAAGAGACATTCATCTTCTTGAATCTTCATTAATCTTCACGTCCTAAATAATTCTCTCATTATT
CAAGTTGCAAATAGAAAGAGGTATTTCCCAATTAGAAATAAAATATTGGGCTTCAGTTTCAAAAAAAAATCTTGAAAGC
TACATGCTCCATAGACTTCTAAGCAAAGAGAGTCAAGAGACAATCATCTTCATAACTAACACCAATTGTCATAAAGCTT
CCAAATGTAAACCTAACTTTTTTGTAGGAGTAAAAATAAAAAAGCCAAACAAATGAAAAGCTGTGAACCTTTCTAAAC
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GGAAGATTTAAACAATTTAAATATATGCTCTAGTTGTTTGATTTAAATGCTTTTTTGTGCCAAAGAAATTCAGGATAGA
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CCTTTAATTTTCATGCCATAAGTCAGTGCTCTTGACAAAGTATACTCTGAACATATTTATAAATCATTATTTTTTCCAAG
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CATAACAGACCCATCCCTGAGAGCAAGGGAGAGAGAGGAAAGCCCTGGCCTTTCTTTTTGTACCTTGATTTTTTCCATTC
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ATTCTACAATAAAAAACCATTAGGATGCCTATGAGCTTTGTTTTCTGCATAATATCTGAATTACAAATTGTATTTAACA
AGTAAACTAAGTTGTGTCCGACTAATTGAAAACCACCTTTGGTTAATGTTACCTCTTTTTTGTGTCCATTTAAATCCATT
AAATCTTTCTTACTTTTGCCTTTAAATTTAGAGTAATCTATACAATTCATGCTACTGACTCTGCCTTTTAAACACACA
AATATTAATAAAAAAAGAGACATGTTCTGGTGTCTTCCCTCAGCCAGTGAGATGTGTTACACAGTAGTCTTTGTAT
CAGCCTTTGGTTTTCTGATAATTTAGAAAGCTTTCAATCCCATCACTTATGAAAGTCTGTGAACAATAATTTTTTATTTA
AAGATTTCTTACAGTGCCGAATCTTACAACCATTTATAAATTCATGTATGTTTTCTTAAAGTTACAGAACTCTTTTCA
ATTTTTTAAGCCTTAACCTCTTGTCTGCATAGCAAATCCTATTTATTTTTTAAAGCTGAATTCAAGTGTCATCTCTTA
AATGAAGATTCCTGATCATGTCAATCAAAGAGAGCTTTCTCCTCTGAAATCTTTTTTTTTTAAATATTAGTTTCAGGGG
CACATGTTTAAAGTTTGCTCTAATAGATAAATGGTGTGTTTTGGGGGTTTGGTGTACATATTTTCGTCAACCCAGGTAA
TGAGCATAGTACCCAATATGTAGTCTTCTGATCATCACCTTCCTCCTACCCCTCAATAGGCCCCGCTGTCTG
TCCTTCTCTTCGTGTGTCCATGTAACCTCAATGTTTAGCTCCCAATTATAAATGAGAACATGCAGTATTTGGTTTTCTG
TTCCTGTGTTAGTTCTCTTAGGATAATGGCCTCCAGCCGCATACATGTTGCTGCAAAGGACATGATCTCATTTCTTTTT
ATGGTTGCATAGTATTCCATAATGTATATGTACCACATTTTCTTTATTTCAGTCTAATGTTGTTGACCATTTTGGTTGAT
ACCTTGTCTTTGTTATTGTGAATAGAGCTGCAATGAACATATGCATGTGTGTATCATATGGTAGAATGATTTATATTC
CTTTGGGTACATACCCAATAATAGGACTGCTGGGTCAAATGGTGGTTCTGTTTTAAGTTCTTCGAGAAATTGCCAAACT
GCTTTCCACAGTGGCTGGACTAATTTACATTCCCCTAGCAGTGATAAATATTCCTTTTTCTTTGTCAGTATCACCAAC
AACTGTTACTTTTTGACTTCTTAATAATAGCCATTCTGACTGGTGTGAGATGATATCTCATTGTGGTTTTGATTTGCAT
TTCCCTAATGATTAGTGATGATGAGTATTTTTTTCATATGCTTGTGGCCGTGTAAATGTCTTATTTTGAAAGTGTCTT
TGCCCACTTTTTAATGGGCTTGTTTGTGTTTTTGTCTGCAATTTGTCTAAGTTCTTATAGATTCTGGATATTAAACC
TTTGCCAGATGCACAATTTGTAAATATTTTCTCCCATCCTGTAGGTTGTCTATTTACTTTGTTGATAGTTTCTTTGCT
GTGCAGAAGCTCTTTAGTTTAAATTAGGTCCCCTTGTCAATTTTGTGTTTTTGTGCAATTGCTTTTGGCATATTCATCA
TAAATCTTTGCCAGGGCCTATGTTTAGAATGGTATTTCTTAGGTTTTCTTCAATGGTTTTTATAATTTTACATTTTAC
ATTTAAGTGTCTAATCCATCTTGAGTTGATTTTTGTATATGATCTAAGGAAGCTGTCCAGTTTCAGTCTTTGGCATATG
ATTAGCCAGTTGTCCAGAACCATTTATCGAATAGGGAGTCTTTCCCATTTGTTTTTGTCAACTTTGTTGAAGA

Fig. 9.41

TCAGATGGTTGTAAGTGTGTGGGTTTATATCTGGGCTCCCTATTCTGTTCCAGTGGTCTATGTATCTATTTTTGTACCT
ATACCATGCTGTCTTGGTTACTGTAGCATTGAAGTATAGTTTGAAGTTAAGTGATGTGATTCCCTCCAGCTTTATTCTTT
TTTGCTTAGAATTGCTTTGGCCATTTGGGCTCTTTTTGGGTTGCATATGAATTTTAGAATAGTTTTTCTAGTTCTGT
AAGAATATCATTGTTTCATTTGACAGGAATAGCATTGCATATGTAAATTGATAAATTCCTGAAAACATTCAACCTCTCGA
GACTGAACCAGGAAGAAATTTAAACCCTGATCAGACCAATAGCAAGTTCCAAATTTGAATCAGTAATAAAAAGGCTACC
AGCCAGAAAAAGCCTTGGACCACACAGATTACAGCAAAATTTCTANCAGACATATAAAGTAGAGCTGGTACCATTCTTA
CTGAAACTATTCCAAAAAATTGAGGAGGAGGAACCTCTCCCTAACTCATTCTATGAGGCCAGCATCATCCTGGCAAAGA
CAAAGCAAAGACATAACAACAACGTAAAAACTTCAGACCAATATCCTTGATGAACATAGATGCAAAAATCCTTAACAAA
ATACTAGCAAACCTGAATCCAGCAGCACATAAAAAAACTAATCCCCCTCTCCCTCTCCCTCTCCCTCTCCCTCTCTCTCT
CCCCACGGTCTCCGTCTCCCTCTCTTTCCACGGTCTCCCTCTGATGCCGAGCTGAAGCTGGACTGTACTGCTGCCACCT
CGGCTCACTGCAACCTCCCTGCCTGATTCTCCTGCCTCAGCCTGCGAGTGCCTGCGATTGCAGGCCTGCGCCGCCACAC
CTGACAGGTTTTTCGTATTTTTTTGGTGGAGACGGGGTTTTCGCTGTGTTGGCCGGGCTGGTCTCCAGCTCCTAACNGCGA
GTGATCTGCCCAGCCTCGGCCTCCCGAGGTGCTGGGATTGCAGATGGAGTCTGGTTCACTCAGTGCTCAGTGGTGCCCA
GGCTGGAGTGCAGCGGTGTGATCTCGGCTCGCTACAACCTCCACCTCCCAGCCGCTGCCTTGGCCTCCCAAAGTGTCTG
AGATTGCAGCCTCTGCCCGGCTGCCACCCCGTCTGGGAAGTGAGGAGCGTCTCTGCCTGGCCGCCCATCGTCTGGGACG
TGAGGAGCCCCCTCTGCCTGGCTGCCCACTCTGGAAAGTGAGGAGCGTCTCTGCCCGGCCNCCATCCCATCTAGGAAGTG
AGGAGCGCCTCTTCCCGACCTCCATCCCATCTAGGAAGTGAGGAGCGTCTCTGCCCGGCCGCCCATCGTCTGAGATGTG
GGGAGAGCCTCTGCCCGGCCGCCCGTCTGGGATGTGAGGAGCGCCTCTACCCGGCCGCGAACCCTGTCTGGGAGGTGAG
GAGCGTCTCTGCCCGGCCGCCCGCATCTGAGAAGTGAGGAGACCTCTGCCTGGCAACCGCCCCGTCTGAGAAGTGAGGA
GCCCCCTCCGCCCGGCAGCTGCCCGTCTGAGAAGTGAGGAGCCCCCTCCGCTGGCAGCCACCCCGTCTGGGAAGTGAGG
AGCGTCTCCGCCCGGCAGCCACCCCTGTCTGGGAGGGAGGTGGGGGGGTGAGCCCCCGGCCGCCAGCCACCCCATCCG
GGAGGGAGGGGCGCCTCTGCCCGGCCGCCCGTCTACTGGGAAGTGAGGAGCCCCCTCTGCCCGGCCACCACCCCATCTGGGA
GGTGTACCCAACAGCTCATTGAGAACGGGCCATGATGACAATGGCGTTTTGTGGAATAGAAAGGGGGGAAAGGTGGGG
AAAAGATTGAGAAATCGGATGGTTGCTGTGTCTGTGTAGAAAGAGGTAGACATGGGAGACTTTTCATTTTGTCTGTAC
TAAGAAAAATTTCTTCTGCCTTGGGATCCTGTTGATCTGTGACCTTACCCCCAACCTGTGCTCTCTGAAACATGTGCTG
TGTCCACTCAGGGTTAAATGGATTAAGGGCGGTGCAAGATGTGCTTTGTAAACAGATGCTTGAAGGCGGCATGCTCGT
TAAGAGTCATCACCCTCCCTAATCTCAAGTACCAGGGACGCAAACTGTGGAAGGCCGAGGGACCTCTGCCTAGG
AAAACCAGAGACCTTTGTTCACTTGTTTATCTGCTGACCTTCCCTCCATTATTGTCTATGACCCTGCCAAATCCCCCT
CTGCGAGAAACACCCAAGAATGATCAATAAAAAAAAAAAAAACAAACAAACAAACAAACAAACAAACAAACAAACAAAG
GTTTGAGCTAAGAACTTGCAGGAGACAAGGAAATTAGTCAAGCAGAAGGATATCTGGGGGAATGGCATGCGAGGCAGAA
GGGAAAGCTAGGGTCGAGGCCCTCAGGGAAAGAAGCAAGGCCAAGGGGCTGGAGTAGAGGGAGGAAGAGGGGAAGTAGT
GGAAGATGAGACTAGCTTTACTACTGATTATGATGTAAGAATAGTGGCCAGTTTCCCTTTCCAACCTTGGGCCCCGCGAGAA
TGGCTCCTGCAAAGAAGGGTGATGAGAAGAAGAAGGGTCAATTCGCCATCAACGAGATGGTGACCCGAGAATATCCCATC
AACATTCATAAGTGCATTCATGGAGTGGGCTTCAAGAAGCGTGCCCTCAGGCACTCAAAGAGCTCCGGAAACTTGCCC
TGAAGGAGATGGGAACCTCAGATGCACACTTTGATACCAGGCTCAACAAAGCTGTCTGGGCCAAAGGAATAAGCAACGT
CTCATACTGTATCCATGTTCCGTTGTCCAGAAAATGTAATGAAGATAAAGATTTACCAAACAAGCTCTATACTTTGGTT
GCCTACGTACCTGTTACCACTTTAAAAAAATCTACAGTCGGTGTGAATGTGAACCTAATGCTAATCATCAATATACCA
AATAAAGTTATAAAATTGTTTAAAAAAACAAAAAAACAAAAAAACAAAAAAACAAAAAAACAAAAAAACAAAAAA
AGGATGCAAGACTGATTTCAACATATGCAATCAATAAATGTGATTCACCACATGAACGGAATAAAAAACAAAAACACA
TGATCATCTCAATAGATGCAGAAAAGGCTTTTGATAAAATTCAGCAACCTTCATGTTAAAAACCTTCAAAAAACTAGGC
ATTGAAGGACATACCTCAAAATAGTAAGCCATCCACAACAAACCCACAGCCAACATCACACTGAATGGGCAAAAGCTG
GGAGTATTCCTCTTGAGAACTGGAACAAGACAAGGATGCCCACTCTCACTGCTCCTATTTAACATAGTATTGGAAGTCC
TAGCCAGAGCAATAAGGGAAGAGAAAGAAATAAGAGGCATCCAAATAGGAAGAGAGAAATCAAACCTACCTCTGTCAA
CTACCTAGAAAACCCCATAGTTTTTGCCCAAAGCTCCTAGATAACTTCAGCAAAGTTTCTGGATACAAAATCAGTAGC
ATTTCTCTACACCGATAATGTCCAAGCTGAGTGCCAAATCAAGAGCATAATCCTATTCACAATAGCCACAAAAATAAC
ATATCTAGAAATATAGCTAACCAGTGAAGTGAAAGAGCTCTCCAATGAGAATTACAAAACACTGCTGAAAGAAATCAGA
GATGACATAAACAATGAAAAACATTCCATGCTCATGGATAGGAAGAATCTCCTCTGAAATCTTATAGCTAGAGAAAC
CATAACATTTATCATACAAATTTGGCATTTTTTTGAGGGTGAATGAATGGGAGAACTATTAGTAGTCACATTGGGACAAT
AATCATAAACTGGAACCTATCCTGTGTAAACTGGATGTATTGTCAACCAAGTTATTGCCTTTTGGGATCCTAACTGTTAG
TATCAAGGACTTTGGACCGAAGTCTGTCTGGTTCCAAATCTGGTTCTCAGCATGACTTTGAGTAGATTATTATATCCT
TTGCCATTCACTTATTCTTTTTATAGCTTCTTACTAAGCTTGTATTAGATGTGTGCAAATTGTAAGGCATTGGTAACAC
AATGGTAAATAATATTTATGGTTTGTGCTGTATGAAGCTTACATCCAACCTTTATCTACAAAGTGGGGCCAAGGATACC
TACATTATGAAAATTTCTTTATATCATCGTGAGAAGTAAATACTATTTGTGAAGCATTTAGTAGGATTTTCAGTATATAG
GAAGTCTCAGTGATAACTATAATTTTATTTGTATTTTTTCATATGCCATTTTGTGTCATAGTCGCATGTATAATTGTAA
GGTTTTTGAGAGTAGGAGGCAAGGCTTATGAACATTTACATACCTTCAGGCATCAGTATAGTAGCCTCTCATTGCATGA
ATGAATTTATTGAGTGAATTCCACTAAACACCAGGTATCAAACCTTTTTTTTTTCTCATTATCTCTTCTGTTTTTTGTT
CCTCTGTTGTTTCTTTCTGCTCTTTTTCAGAGCATTGTAATATTTTTTAGTGTGCCATTTTATTTTTTCTATTGGCTTT
TAGTTATACCCCTTATCTCTCTGTTTATCATTTATCTTCTTTTAAATTTAATTTTCTTGATTGCTCAAGGGCTAAAA
ATATGCAGCCCTATGCATTAGAATCTATTCTACTCAGTATTGTACCTCTCCACATTTCACACTTAACATTTGACAAA
TGTAGTAAACTTACAACATTTTATTTCAATTTAGTCATCCTCTTTCTTTTATTTTCTTTTTTATTTTTTTGAGACAGGGT
CTTGCTCTCACCAGGCTGGAGTGCAATGGCACAATCATGGCTTACTGTAGCCTCAAACCTCTGGGGCTCAAGCAGTCCT

Fig. 9.42

CCTGCCTCGGCCTCCTGAGTAGCTGGGACTCAGTCATGTGCCACCACACCTAGTTAATTTTTTATTTTTATTTTTTAGTA
GACATGAAGTCTCACTATATTGCCCAGGCTGGTCTCAAACCTCCTGGGCTCAAATGATCCTCCCTGCTCTGCCTCCCAA
GTGCTGGGATCACAGGCATGAGCCACAACACCTGGCCTCATCTCCCTTTCTTTGTATGGTTATTGTCATATTCAATTATT
TTAACATGTCATAACCCCAACCTTAGAGTGTTTTCACTTTTGCCCTTAAATAGTAGTCTTTAAGGAAATTCTAAGAAAGA
AAACATACTACTTTTATATTTAGTAATTTACTTAAACATTTCCAGTGCTCTTTTTCTTTCTTACTGTAGATCTGAATTTCC
ATCTGGTATCAGTTATCTTCAACTTTTTGTATTTCTTATAGTGAGGTCTGCTGGTGATTAATTTTCTCAGCTTTTATT
TATCTGAAAGTGCCTTTATTTTTCTTCCCATTTTTGAAGAATATTTTTGTGTGGTTGTAGAGTTCTGAGTTAACAGCTTTG
TTTTTCGTTTTTCCFTTCAGCGATTTAAAAATTCCATGCCATTGGGGTTTTTCAGTCCCCACTGTTTCTCATAAGCCCATG
ATCATTCTTATCCTTGTTCAACTGTACGTAATGTGTGTCTTCCCTCTGGATGCCCTTAATATTTCCCTCTTTATCTTGGA
TGTGCTGCAGTTTTATTACAATGTGCCTGGGTGTGGTTTTCTTTATATTTATTCTTCTTGGGGTTTTGCTAAACTTCTTG
GATCTGTTTGATCGTTTTTGAAAAGTTTTTCAGCTGTTATTCTTCAAATATTTTTTTCTGGCCCATTTCTCACTCGCTTTT
CTTTCTAGGATTCCAATTACATTTACGTTAAGACTATTTGGTATTGTCCTACCTATTCCTGAATCTTAGTTCATCTTCC
TGATTATTTTCCCCACTCTTCTTTCTAAGATTAAATAATTTCTACTGATCTGTTTTAGGTTCTCTTTTCTTCTGCCAT
CTCTAATCAGCTATTCAGACCTGTCCACTAAATTGTTCATTTTTATTTTTTTTTTAGTTCTAGAATTTTCATTTGGTTAT
TTTCTCTGCCTAGACTCCTCTATTCACTCCTTGAGATCATATTTCTGTCAATTCTTTGGACATATATATTTAAATTTCT
TTAACATATTTACAAAAGCTGCTTTAGATTCTTTGTTTGACATATCTGGGTAATTTTGAGATTTGTTTGTATTTACTG
CTTTTTTGTCTGCCTATGTATCATATTTTCATTTTTTTTAAATTATGCTTTAAGTTCTAGGGTACATGCGCACAGCGTGC
AGGTTTGTACATATGTATACATGTGCCATGTTGGTTTGCTTCACCCATCAACTCATCATTTACATTAGGTATCTCTCC
TAATGCTATCCATCCCCCACCACCCCAACAGGCCCTGGTGTGTGATGTTCCGCCACCCTGTGTCCAAGTGT
TCTCATTGTTCAATFCCCACCTATGAGTGAGAACATGTGGTATTTGGTTTTCTGTCAATTGTGATAGTTTGCTGAGAATG
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ATATGTGCCACATTTTCTTAATCCAGCCTATCAATGATGGACATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATAG
TGCCGCAATAGACATACGTGTGCATGTCTTTATAGTAGAATGATTTATAATCCTTTGGGTATATACCCAGTAATGGGAT
GGCTGGGTCAAATGGTATTTCTAGTTCCAGATCCTTGAGGAAACGCCACACTGTCTTCCACAATGGTTGAACATAATTA
CACTCCCACCAAGTGFAAAGGCATTCCTAFTTCTCCACATCCTCTCCAGCATCTGTFGTTCCTGACTTTTTFAATAATT
GCCATTCTAACTGGTGTGAGATGATATCTCATTGTGGTTTTGATTTGCATTTCTCTGATGACCAGTGATGATGAGCATT
TTTTCATGTGTCTGTTGGCTGCATAAATGTCTTCTTTTGAGAAGTGTCTGTTCATATCCTTCGCCCACTTTTTGATGGG
GTTGTTTGTTTTTTTCTTGTAACCTTGTTTAAAGTTCTTTGTAGATTCTGGATACTAGCACTTTATCAGATGGGTAGATT
GCAAAAATTTTCTCCCATTTCTGTAGGTTGCCTGTTCACTCTGATGGTAGTTTCTTTTGCCATGCCGAAGCTCTTTAGTT
TGATTAGATACTATTTGTCTATGTTGGCTTTTGTGTCATTGCTTTTTTGGTGTTTTAGTCATGAAGTCCTTGCCCATGC
CTGTGTCTGAATGGTATTTGCCTAGGTTTTCTTCTAGGGTTTTTATGGTTTTATGTCTAACATTTAAGTCTTTAATCCA
TCTTGAATTAATTTTTGTATAAGGTGTAAGGAAGGGATCCAGTTTCAGCTTTCTACATATGGCTAGCCAGTTTTCCAG
CACCATTTATTAATAGGGAATCCTTTCCCCATTTCTTGTTTTTGTGTCAGGTTTGTCAAAGACCAGATGGTTGTAGATGT
GTGGTGTATTCTGAGGCCTCTGTTCTATTTCTTGGTCTATATCTCTGTTTTTGGTACCAGTACCATGCTGTTTTGGTT
ACTGTAGCCTTGTAGTATAGTTTGAAGTCAGGTAGCGTGATGCTTCCAGCTTTCTTCTGTGCTTAGGATTGCTTTGG
CAACGTGGGCTCTTTTTTGGTTCCATATGAACTTTAAAGTAGTTTTTCCATTTCTGTGAAGAAAGTCATCGGTGGCTT
GATGGGGATGGCATTGAATCTATAAATTACCTTGGGCAGTGTTGGCCGTTTTTCACAATATTGATTCTTCCCATCCATAAG
CATGGAACGTTCTTCCATTTGTTTGTGTCTCTTTTATTTTCGTTGAGCAATGGTTTGTAGTTCTTCTTGAAGAGGTCCT
TCACATCCCTTGTAAGTTGGATTCTAGGTATTTTATTCTCTTTGTAGCAATTGTGAATGAGAGTFCATCATGATTTG
GCTCTCTATTTGTCTGTTATTTGTATATCAGAATGCTTGTGATTTTGCACATTGATTTTGTATCCTGAGACTTCGCTG
AAGTTGCTTATCAGCTTAAGGAGATTTTGGGCTGAGATGATGGGGTTTTCTAAATATACAATCATGTCTGCAAACA
GGGACAGTTTGACTTCCTCTTTTGCTAATTGAATTCCTTTTATTTCTTTCTTGCCTGATTGCCCCAGCCAGAACTTC
CAACACTAAGTTGCATAGGAGTGGTGAGAGAGGGCATCCTTGCTTGTGCTGGTTTTCAAAGGGAATGCTTCTAGTTTT
TGCCCATTCAGTATGATATTGGCTGTGGGTTTTGTCAAAAATTGCTCTTACTATTTGGAGATACATTCCATCATTATGTA
GTTTATTGAGAGTTTTTAGCATGAAGGGCTATCGAATTTTGTGTAAGGCCTTTTCTGCATCTATTGAGATAATCATGTG
GTTTTTGTCAATTGGTTCTGTTGACGTGATGGATTATGTTTATTGATTTGAGCATGTTGAACCAGCCTTGCATCCCTGGG
ATGAAGCTGACTTGATTGTGGCAGATAAATTTTTGATGTGCTGCTGGTTTCAGTTTGCCAGTATTTTATTGAGGATTT
TCTCATCGATGTTTCATCAGGGATAATGGTCTAAAATTCTCTTTTTTTGTTGTGTCTCTGCCAGGCTTTGGTATCAGAAT
GATGCTCATAAATGAGTTAGGGAGGATTCCTCTTTTTCTTTTGTATTGGAATAGTTTCAGAAGGAATGTTACCAGCTC
CTCTTTGTACCTCTGGTAGAATTCAGCTGTGAATCCATCGGGTCTGGACTTTTTTTGGTTGATAGGCTATTAATTATT
GCCTCAATTTCAAGAGCCTGTTATTGATCTATTCAGGAATTCAACTTCTTCCCTGGTTTTATTCTTGGGAGGGTGTATGTGT
CCAGGAATTTATCCATTTCTTCTAGATTTTCTAGTTTATTTGTGTAGAGGTGTTTATAGTATTCTCTGATGGTAGTTTG
TATTTCTGTGGATCAGTGGTGATATCACCTTTATCATTTTTTTATTGCATCTATTGATTCTTCTCTTTTTTTATTAGT
CTTGCTAGCAGTCTATTTTTTTTGTATCTTTTTCAAAAACCAACTTCTGGATTCATTGATTTTTTTGAAGGGTTTTTTGTGT
GTCTCTATCTCCTTCAGTTCTCCTCTGATCTTAGTTATTTCTTGTCTTCTGCTAGCTTTTGAATTTGTTTGTCTTGTCT
TCTCTAGTTCTTTTAAATTGTGATGTTAGGGTGTGATTTTAGATCTTTCCTGCTTTCTCTTGTGTGCATTTAGTGCTAT
AAATTTCCCTCTACACAGTGCTTTAAATGTGTCCAGAGATTCTGGTATGTTGTGTCTTTGTTCTCGTTGGTTTCAAAG
AACATCTTTATTTCTGCCTTCATTTCAATTATGTACCCAGTAGTCATTCAGGAGTAGGTTGTTTCAGTTTCCATTTAGTTG
AGCAGTTTTGAGTGAGTTTCTTAATCCTGAGTTCTAATTTGATTGCATTGTGGTCTGAGAGACAGTTTTTTTGTAAATTC
TGTTCTTTTACATTTGCTGAGTAGTGCTTTACTTCCAATTATGTGGTCAATTTTAGAATAAGTGTGATGTGGTGTCTGAG
AAGAATGTATATTCTGTAGATTTGGGGTGGAGAGTTCTGTAGATGTCTATTAGGCCCGCTTGTGCGGAGCTGAGTTCA

Fig. 9.43

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AGTCCTGGATTTTCCTTGTTAACCTTCTCTCTGGTTGATCGGTCTAATATTGACAATGGGGTGTTAAAGTGTCCTATTAT
TATTATGTGGGAATCTAAGTCTCTTTTATAGGTCTCTAAGGACTTGCTTTATGAATCTGGATGCTCCTGTATTGGGTACA
TATATATTTAGGTTAGTTAGATCTTCTTGTTGAATTATCTCTTTACCATTTATGTAATGCCTTTCTTTGTCTCTTCTGAT
CTTTGTTGGCTTAAAGTCTGTTTTGTCTAGAGACCAGGATTGCAACCCCTGCTTCTTTTTGTCTTTCCATTTGCTTGGTAG
ATCTTCTTACATCCCTTTATTTTGAACCTATATGTTTCTCTGCATGTGAGATGGGTCTCCTGAATACAGCACACTGATG
AGTCTTGACTCTTTATCCAATTTGCCAGTCTGTGTCTTTTAATTGGAGCATTTAGCCCATTTACATTTAAGGTTAATAT
TGTTATGTGTGAATTTGATCCTGTCTATTATGATGTTAGCTGGTTATTTTGCCCATTAGTTGATGCAGTTTCTTCATGGT
GTCGATGGTCTTTACAATTTGGCATGTTTTTGCAGTGGCTGGTACCAGTTGTTCCCTTTCCATGTTTAGTGCTTCCCTCA
GGAGCTCTTTTAGGGCAGGCCTTGTTGGTGACAAAATCTCTCAGCATTTGCTTGTCTGTAAAGGATTTTATTTCTCCTTC
ACTTACAAAGCTTAGTTTGGCTGGATATGAAATTCTGGGTTGAAAATTTTCTTTAAGAATGTGGAATATTGGCCCC
CACTCTTTTCTGTCTTATAAGGTTTCTGCCGAGAGAGCTGCTGTTTGTCTGATGGGCTTCCCTTTGTGGACAACCCGAC
CTTTCTCTCTGGCTGCCCTTAACATTTTTCTTCTCATTTCAACCTTGGTGAATCTGAAAATTATGTGTCTTGGGGTTGC
TCTTCTTGAGGAGTATCTTTGTGATGTTCTCTGTATTTTCTGAATTTGAATGTTGGTCTGCCTTGCTAGTTTGGGGAAG
TTCTTGTGGATAATATCCTCCAGAGTGTTCCTCAACTTGGTTCCATTCTCCCCGTCACCTTCAGGTACATCAATCAGAC
GTAGATTTGGTCTTTTTCACATAGTCCCATATTTCTTGGAGGCTTTGTTTGTCTTTTACTCTTTTCTCTAAACTT
GTCTTCTTGCTTTATTTTCAATTCATTTGATCTTCAGTCAGTGATATCCTTTCTTCGACTTGATCTAACAGGCTATTGAAG
CTTGTGCATGCATCACGAAGTTCTCGTGCCATGGTTTTTCAGCTCCATCAGGTCATTTAAAGTCTTCTCTACACTGTTTA
TTCTAATTTGCCATTCGTCCAACATTTTTTAAAGATTTTTCAGCTTCCCTTCAATGGGTTAGAATGCCCCCTTTAGCTTG
GAGAAGTTTGTATTATTACCATCCTTCTGAAGCCTACTTCTGTCAACTTGTCAAAGTCATTCTCCGTCCAGCTTTGTTCCA
TAGCTCGTGAGGAGCTGTGATCCTTTGGAAGAGAAGAGGCACCTCTGGTTTTTAGAATTTTAAATTTTCTGCACCTGGTT
TCTCTCCATCTTTTTTGGTTTTATCAACCTTTGGTCTTTTCATGTTGGTGACCTACAGATGGGGTTTTGGTGTGGATGTCC
TTTTTGTGATGTTGATGCTAFTCTTTCTGTTTGTAGTTTTCTCCTAACAGTCAGGTCCCTCAGCTGCAGGTCTGT
TGGATTTTGTCTGGAGGTCCACTCCAGATCCTATTTGCCTGGGTATCACCAGTGGAGGCTACAGAACAGCAAATATTGCT
GCCTGATCCTTCTCCATAAGCTTCGTGCTAGAGGGGCACCTGCCTGTGTGAGATGTCTGTGAGCCCTACTGGGAGGT
GTCTCCCAGTTAAGCTATGTGGGGGTCAGGGACCCACTTGAGGAGGCAGTCTGTCCGTTCTCAAAGCTCAAACGCCATG
CTGGGAGAACCCTGCTCTCTTCAGAGCTGTCAGACAGGGATGTTTAAAGTCTGCAGAAGCTTCTGCTGCCTTTTTTTCA
GCTATGCCCTGCCACAGAGGTGGAGTCTATAGAGGCAGTAGGCTTTGCTGAGCTGTGGTGGGCTCTGCCAGTTCCAG
CTTCTGGCCACTTTGTTTACCTACTTAAGCCTCAGCAATGGTGAACACCCCTCCCCCACCAGGCTGCTGCCTCGCAG
GTCAATCTCAGATTGCTCTGCTAGCAATGAGCAAGTCTCCATGGGTGTGGGACCTGCTAAGCCAGGCACAGGAGAGAAT
CTCCTGGTCTGCTGGTTGCTAAGACAGTGGGAAAAGCGCAGTATTTGGGCGGGAGTGTCTGTTTGTGATGGCTTCCCT
TTGCTAGAAAAGGGAAATCCCCAACCCTTGTACTTCCCAGGTGAGGCGATGCCCCACCCTGCGCTGGCTCCCGCTCC
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TAGTCCAAATCTTTGAGAAATCCCAGGTGCTTTCCCAACCTATTCAACCTGGCAGTATTCAGTGTGATAGAGGATGTT
TTTCTTGATGATAGGCTTTGTTTTAGACTTTACTGGAGTCATAGGACTTACTTTAGGACATAGTCTTTACTTGTAGAGA
GGTACCAACTTTCTGTTTCTCAGGTAGATCCCAGGGGTGTCAAAGTAGTATTTATTTATGAGCTCTCTCAAACCCATAG
GACCTGAACCTGCAATGATGTCTAGTACTATTCTTCTTCCAGCATTAATTGACCTCCACTATTTCTGTTCTCTCAACCTG
ATAACATTTTCTCTCTGTTAAGCCTCCAGTATTCTCACTCTGCAAATGTATGGTGGTGTCTCAGTCACAGATTTGTCC
CATGTCTGGGACAAATCTCTGCAAACTTCTGAGACTTCTCTGTGTTAAAGTCTTTACTCTCTAAGACTCTGCTTTATA
GATGCCAGCCATGCCAGCTGCCTCAGACTCCAGCTCTTTTTGTGATGTTTAGGAAAATATCCTTAFTTACAGAGGTGGA
CAFTCGTGGGCAGAGGGATTTTCAAGTCTGGATTGGCTTGTAGCCACTGTTTGAAAACGGTTTCCCTCATATATTTTACT
TAGTTTTGTAAAGTATTTTCTGTGAGACAGATAATCTGTTACTAGTTACTCTATCATAGCTGGAAGCAGAAATATATAGG
TATCAATTTGATTTGCAATTGTTTCTAGTTTACAATGCATTCTGCCTATCTTAAAAAATTTGTAATTTCTAATCATTTTA
TTTTTGATCAGGGAATGTATTTATTTGATTACATGAATAAAATTCAAAGGCTATAGAAGAATATGTAGCAAAAAATCTC
TCTTCTAATCTAGTAGCCTAATTCTTCTCCTCAGAGGTGATAGATATTACCAGTTCTTTTTGTTCCCTTTTCCAGACATAT
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TTTTTGGGCTACATGATACTCAGTTTTATGAAATACCATGAATTATCTCACCAGATTCCCTATTTTTTACTCATACTTTTT
GGTTATTATAAACAATGCTTCATGAACACTTTTGGGTAAACCATTTGGTATATGCCAGTATATCTGTAGGATACATTCA
TAAAAGTGCAATTGCTACCTGAAAATGTATGCACATTTGTTACTTAGTTAAATGTTGCCAAATTTCCCTCCAGCAGCTG
TGTTAGTGGCTGTAGCCTCGTTAAAAATGTATGGGAGGAATGCTGACATTTTTGTTTTTACCTGCAGTATCCATTTTCTC
CATGGTACATCTACAAAATTGGCTTCTATTTTTTATTTCTTATGTTTATCTTAATCTTTTAATAATCTTTCCCTGATTAA
CTTACAAGTTTCTTGTATATAACAAATAAGGGGGCTAAGGGGGGAGGGGGATGGTTAATGAGGAAAAAATAGAAAGAATGA
ATAAGACCTACTATTGATAGCACAACAGGGTGACTAGAGTTAGTAACTTATTTCTACAATTAAAAATAACCGAGAGTG
TAACTGGATTGTTTGTAAACACAAGTGATAAATGCTTGAGGGAATGGATACTCCATTCTCTATGATGTGATTATTTTACA
TTGCATGTCTCTGTGTAAACATCTCATGTACCCCATAAATATATTTATACCTACTATGTACCCAGAACAAATTAATAA
AAAAATTTAAAGTACAAAAAAGACAAAAAGAGTCACACAGAAAAATAAAGGAAAAATAGTCTATAGAAGGATAAAAAACA
AAACAAAAAACAATAAGATAAGCAGCAGCGATTTATATAAATTAGAGTTATTATTACTGTGGTTGCTGTGGTTG
ATTTTCTAATATTCCAGGGCACGAATCAGCTGATAATATCCATTTTAAAGTGCAATTTTATTTCCATTTTAGTCAGT

Fig. 9.44

CTGATTTTCAAATTATTCTCCCCTTTTGTGCTTTAAAAATAGGGGTATAACATATGCTAGCCTGTTCCCAATTTGAAC
TCGTGTCTTTCAAATGCACAGTCTAACTTTATTAATTCTGTATGGAATCATGTCTGTGAGCTTTCTAGCTTTTCTTTC
TTTCTTGTTTTTTTTTCTTTTACCTCACTTGCTCTTCTGAATTTGAGGAGCTAAAGATATTGTGACAAGACTTAGCTAC
TCAATGAACATTCCGAATCAATGTTTTCTGTGGAACAGTGACTCTGATGGAATCTTCTCCACTTCCTGGAACATTAA
ATTGTCACTAGGTAAATTCAGGAGTCTCTTTTTTCAGGTTCTTGTCTGCAGTATACTGAAAAGTGAATTGTGACCTCATTG
TTATGAAATTTGGAAGTGTTTATCTTTAATGTTAGCTTAGAGTGTTTTTTTTTTCCCTCTCTCTTGCAATCCCTTTAAAC
CCTGACCATTAAACATTATAGACAAGGACATTTCTCATGTGTTTGTCTTTCTCTAAATCTGTTAGTACTTTAAACTGT
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CAGAACAAGATATGCAGAAGTCTGGTTTTTGGCCCTCCAGGAGTGCAATGGTTGTGAGTGTAGGATCTGAAGTCAAGCA
GACAGATGTAGGCTGGATGATCTTGGGTGAGGTACACTAAATCTCAGTTCTCTCATCCGTGATACAATTGTACTCATCT
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ATGTGGTCTTTTCAATTGCTGTCAATCAGTTTTTAAAATCTGGTTGCCCAAACACCATTGTGGCTATTACCATGACTAC
TTCATGTTCTGGCTGTTCTGGCAGTACAAATGAGTCTGGACATCTACTTCAGGTAAATTGGATGTTTGATTAATTAATA
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ACATCTCAATGGCTTTATATATTACATATTTATTTCTTGTTCTGATCACATTACAATGTGGATTGGGGGTAAAGTGGGA
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CATTTACAAACCTTGGGCTAGATACAGAGTGTGATTGGTGCATTTACAAACCCTGAGATAGACACAGGGTGTGATTG
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CAAGTCCCCACCAGACTCAGGAGCCAGATGGCTTCACCCAGTGGGTCCCGCACCGGGCGGCAGGTGGAGATTCTGCC
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AGCCTGATCAGATGCAGAGGGTCTAGGAAATAGGTGCTGAAAGAAGGGAACCCATGCACATTAATGAGTACCATTGATT
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CTTTCACCAGATTGAAGTTTCTAAGCTGTTTTTCCAAAAAATAAAGGATAAGGCTTTGGCCCTTTTAGAATCTGGTT
TTATGAAGGTGGAAGGAATGACAAGGCTTAGTGCAAAATATTAAAGTACCTTATGCCCTTTTACCAAGTGTGCACCAC

Fig. 9.45

Fig. 9.46

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CCTTTGAAGGAAGACATCATGCTCGTATTTCTTTACACTTTTTTTTGATATTTAGCTGCTCAATACAAACATGTTGGTG
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TTTAGGGCCACACCCAAGTGGAACTACTTATTTTCAGGATTGGAGCCTCCACAGAGAAAGATCAGAGGTGGGAATGAAA
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TACGAGTTCCAGTAAAGAAGACGGAAGCTACACTGCTTTTCATGTCCAACCTTGGAAATCACACGGCCTCACTTCCACT
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CTCCAGATAAATGAATAGAGAGAGAGACAAAAAAAAAAAAAAAAAAGAATTGGCTGACTCGGGGTGCTGGGTACACAA
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TCTGAGGAGAATCGCTCAGTTTTTAAACAAGTGCCAGTTTACACTAAAATGCTTGTCAATTTATATAAAATATTGGCATT
TGTTGCAATATTGCAAACCAAGAGTCATTGAGTTCCAACTAAACAAGTTTTATGAAACAGTTTGTCTGTCCAAC

Fig. 9.47

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CTTAAAAAGAGAAGGCATTTACTATATGCCAAGCATTGTGCTGCCTGCTTCACGTGCATTTTCTAATGTGGGTTTCACA
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CCTGCCGAGCCTCCTAAGACAGCACTGTGGAGTGGAAACACCTCTGATCTGGGAGCTGGATATTCTGGCCTCCAGACCT
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CTGTAAATGGTTTTGGTTTTCTGGTTTTATAAGTGGGTTACCAGGCAAGTGAGTTTTTCATGTTGAGGCATGGGGCAATGCC
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AGCCCTCAGTTTTTAATTCTATAAATTAATGGATTTTATAGTAGGATGTTAGGACAAGAAATTTATATGTATTACATA
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GGCTGGAGTGCAGTGGTGAGATCTTGGCTCACTGCAAGCTCTGCCTTCCGGGTTACGCCATTCTCCTGCCTCAGCCTC
CCGAGTAGCTGGGACTACAGGCGCCTGCCACCACGCTGGCTAATTTTTTGTATTTTTTAGTAGAGACAGGGTTTCACCA
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ATTAATAGTCTGTATTTCAAGTAGGAGAAGAAAATATAAAGAGATATCCTTCAGTTCCCTTTTCTTTATAGCTTTCTTGA
AGTTTCTATAAGCAATACTAAGAGGTAATATAGCCTCTGTACACCCCCACCCCAAGAAATTGGATTGGAGAAGTGAAC
ACACCTTCTGAGAGTGCAACTATAAAATAGGCCTAGATGGATCAGATGAAAGGAAATAAAAAGCAATTATTTAAGTGG

Fig. 9.48

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GGCTTAGATGTTGCTTAAACTTGGCTTTTCCTCCTCTATAAATAATCAAATTATAGATCTCCAGGGACCTAAAATGTTA
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TGTAATCCCATATTAAGTATTTAGTAAATAGCTATTCTCATTCTAATTCCTGTTATAGGAGAATAGAGGAGAAGTTGG
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TCAGGATCATCTATTTTAAACATATCTGCTCCTTTGTGGGAAGAGTGAGTGTGAAAATTTGGGAACATGCGCAGTGGATA
AAGCTGTTATTTTATGCAAAACAGAAAAGAGCAGTGCCCCCTTTGGGTGGATGTAGCCCTGTACCAGTGCACACAGCTT
GGTATTCTCAAAGAAGCCACAAAACAACCAAGGAGGCTGTTCCAGCTGGGCACCTAGTAGCTGATTCTCTTGGTAACCG
TGGGCTGGGTTTTAGCTNTCACATGCCTGCTTGAGCGGAGTACCTTTGTGTGGTAAACAACCTGGACCCTCTACATGG
TCCCCTGGTGGTGGTAATGGGCAGGCTTAAGCGGTGTGATATTGGTTAGGGGGCTCAAACAACCTCATGGAATGGGCACC
TAAGCAAGGTAGAGAGATGCAGAGCGGGGAGGGAGGAGTTCTGAAGGGGAGTTTGGCTTGTGTTGCTTTTTTTTTTTTC
TTCTATCTTGGCACCCCTTTATTAACAGAACACAGTGTAGATTATGCATTGTATTTTGAAGAAAGAGAGAAATAAAGG
CGTAATCTTTGCTTTTGAGATGTTTTTCAGTCTAACTCAGCAATTCCTTTCCCTCATCCTTTCTGCCTCACTACTTATTA
CATTTACTTCTAGTTCTTCTCTTTAAATTGAAAAAGCAGAGGCAAACACAGACTGTGATAATGGAATTTTATGTTG
TGGTCTTTGATTACACTTCCCTTCAAAGCTTCAATATTTACATTTGTTAATAATAATGTTCTTTGTAGGAAAAATCC
TTAATTTGAAAATGTTGCAGATGGCCTTTTATCAGGAGCTATGGACTGTCAAATAAATAGATATAATTATGAACCATT
GTTGTTGAATGGGCCATCTTAAATAATAGCAGTCAAGTTTCCAGAAATTAACAAAAAACCCTTACATTTTAAAAAGAC
CTCAGAATAATTAGTGGTCTCTGGTTGATTGGTTTTCAACAAGTTATTCAGCAGATGGAGGCTGTCAGATTTCTGCTGG
TACGATTATTTTTGTAACCTCAGTGGTGGAGGGGGTGGTTTTATGCAACTCAGCATCACCTCTTTGTTGAATTGGATTG
GTTTAAGAGAATCCCACCGCACCTGTACATTAAGATGGAGCATGAGGAAGGTCTTGGGACCGAGCCATTTGAAGAAAAT
CTAGGCTCNGGTGGGTCACTTTAGGAGTATTAGTATAGCCTTAGACATTAGCTCTTGGAACTACCCTGAAGGCGAAAAT
GTGAGACAGCTAATATTTCTCTGCAAGAATTCCTTGGGTGCTGAGTTTTGGTTCTTGCCATCAAGAAAATGTTTGTAC
CGGAGCAAGATGTGGAACCCAGATGAGAGGATATTAATGTAGATTCATGAGCTCTGCCAAAGTAATGTCATTACTG
CTGCTCCATCCCTGAAGAAAGATCTTAAACATATGTAAATAGACAAGACAAGTTATAAATTGTAATTTAGTATCTGGTA
ACTGAAAGTCTTTACTTCAATTTGTACTGAGTGATTACCTGAATTTACTAAGGAAAATTTTGGAGGTCACAGATTTGAGT
TGAAGTCAATAAAATAGGATAAAAGTCTAGATGATGAACTTAGCTTTTGTGATTAGAGTTCTGTTTAGCTCTTAAAA
CTGCAGTAAATAAAATGTTATATTAGTGGAAAATACAAATGGATTCAGAAAATATAGACAAATCGATAGGCAATTGAAA
TGTATACATTTTATTTTCAACATATCAAACCTCGAGTTCAAAGTTCTTCATAAAAAACACAATTCCTAAATTAACCTTAC
AATAACTGTGAGCATTCCTTATCCCATTTTCAAGTGTCTACAAACCCAAAGTTATGGGGAGAAAACCTTAAAGGAGGCAAGA
GCTGCCACTATAATTTAAATATATTGTTCTCCACTCTTTTTACATATTCTTGAAAGCAGTTTCAATTAACGGTGACCTTG

Fig. 9.49

TGTAGGAAAAATTAGCATTGTGTCACAAAAATTCTTTGTATGTGTTAGTTTGTGTGCATATTTTGGAGTCTTCATGTTA
AAAGTATAGGACAGACCTACTTGACAAAGGTGAATTTTGTGCGAAATTTTGGGGAGAATATAGATTTGAATTCATGTAAA
TAAGATTGAATAAAAAATCCAGATGACTGAAACATATTTTCATCTTTTGCTAATTAGAGCTTTATTGAGCACTTAAGCTGC
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ACTCAGGAAAAACATTCAATGAAACCTCTTTTTCAACACATAGATTAGTACCCTAAGAGAGTTAAATTAGTGTTTTACTG
TGAAACTATTTTCTTCCACACCAGAATCTATTGTATGTTGGCAAAGAAAAATGGACTAGCCATTCATTATCACCAAGAG
TTTATTACTTACTCTGGCCACCTCAGTGTTTCAAGTGGGTACCTAAAGTGAAAAGAGTTTTAAGTGCTACCTTTGCAGTAT
TAAGTTTGAGTGAATCCAGGGATATTTGCTGTGAGTGTGGCAAATGATTTATTGTGGAATTGGTTGTTGAATTTGAATT
TTTTGTGTGCTGGCAAATTCACATTGCCAAGTCTCTCTTTTAAATTGAAGAGTATTTTAAAGTGAAATGCAATACAAACA
TATCTTATTCTTGAACCTTCTTGATTAAATTGGTAGTGTTTTTTTATTGTAGTAGGCTACTTATAGAGTTCTTTTTTCAT
AAAACCTGAAGGTTCTTCTTCATAAATCTTGACGCTTCAAGGGGAAAAAAGGAAAAAAGAACTTATTTCTCTTAGA
CCCTTACTGTCACACCTCAAGTATTGGACACAATGTGCGATGCTTAAAGACTCTCTTGGCTTAGAAAAAGCCTTTCTCT
CTCTTGGCAAATCAGGCAATGTGAAATCAGTAAGGGCCAGTTCTCACTGTTTTCTTATGAAGTTGTTCCGATGTGTGA
CACATCCACCTTATGGAGGTTTAAAGTTGGTTGTGTCTTGAGGCTCACAAAAGCAGGTTGTGGAATTGGTATAAGCATTT
GATTTATTTGAACATATCTGGTGGTGTGTCTTACTGACCCCTCTGTTATAAGCTTTCTTCAAAAAAGTCCATCAG
AGACTTGAAGTTTCACAATCAAAGCTTTGTTTACTGGGCATATTTTTTCAGCCTCAAGGAAATCTTCCCTGGCTCCCTGT
AGAAGCAATAGTTACACCTTCTTAGGGGAAGAACCCGTTTAAATGTGTTGCACATCTTACCTCTTTTACTTGAATAA
GAAGTTTACAATCTTCCCTGTGGTATCCTTCTTAGGGGAACCTTAGGAAGAAATGTTTGATCATGAGAGTGGTAACTGG
AAGCATGAACTGGAGTACGGTGTTCCTTCTTAAAGAAATTTTGTCTTAAATTTTTTTTTTAAATGGCTTTCCAGAGGTCA
GGCAAAGGCCAGTTTCCCTTGTATATTGGGAATTTTCAAGGTGGGATTAACAAAAGGCTAGAAAATGAAGATGGAAAGGG
ATTTATAGCTACATAGCAGCAGCATGGACAATTTCTTAATGTGCATCTCTAATTAAATATTGGTGTATTAGCTTATTAA
TTACTTCAACAAATATTCATTGAGTATCTCTGCCAGGCAAGAATCTGGGCACCAGAGTTACAAAAGTGAATGAGGCACA
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CATTGCCAATTTTCAAGTTTGTAAATTTGTTCAAACCTTGCAAGTGATAATGTGGTACTTTATTTTGAAGGAGCAAAACAATAA
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GAAAGTTCTATAAAAAGATTTCTCCACAATTCATTTTATAATAGTCTGTGAGTGGTTTATTAACCTACTTTTCTG
TTGAAAATATGGAGTCTCAGTAATTTGTGATTGAGTTAGTCATCCCAGGTTATAAAACCAACAGTTTCAGAAGGGTCTTG
AATGAAATATCTTCTGCCTTTCAATCCAGGGAAGTTTGTACTATACCTCAACAAGGACTAAAGGTAAGGGCTAAACATG
GAACATCACAGCTATCAAATTGATGAGAGAGGCCTTATCACTTCTTAGGTTGCTCAAGAGGATGATGTGAGCAGAATAA
TGGCCCCCAAAGAGGTTTATGTTCTCATCTCTGGAATCTTTGAATACGTTATAGGACAAAGGAGAGTTAAGATTGTGGA
TGGAAATTAAGATTACTAATCAGATGATCTTATAATAGGGAGATTATCCTGGATTATCCTGGTGGGCCCAATTCAATATT
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TTCTTTAAAAATGGAAAAAAGGGCCACAAGCCACAGAATGTGGATGATCTCTAGAGCTGGAAAAGGCGAGGAAAGAGAA
TTTCCCCTAGAGCATCCAGAAAGAACTCAGCCCTATGAACCTCTTGGTTTTAGCCCACTGAGACTCATAGGACTTCTG
TCCTATGGAACCTGTGGCATGATAAACTTTTGTGTTTATGCCACTAATGGTAATTTGTTAAAGTAGCAGTAGGAACT
AATCTGCGGGGTGGCTGTTAGAGCCTGATTTATAGCTATCTACCTTCAGAGCAGACAGTGTGACTGGGAGCCCCCTGTG
ATCACCATGTGGGGATGTTATAGGTTAGATCAATCATTTATTCAATTATTTATTAATAAACATTTATTGCACATCTACT
TGGTACAAGTCACTGTTCTAGGCCAAACAGTAAACAAGACATAAAAAATCTTGCCTTCATGAAGCTATCATTGTAGAAG
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GATAGGAAGCAGTGTGTGTGTGTAATATGCATGTGTATATGTGTGCATGTTTATATTAGGTTTGTAAATTTTAGATGG
GGCTTCTAGGGCAGGTTCACTGAGAAGGTGATATGTAAATAAAGACCCGAGGACTTCTTTTAGGAAGGAAGAAACAGCC
AGTGACTTTCTATCCCAAGCAAAGGAAATGCAAAGGCTCTGAGGGCAGAGGGTTCTTGGCACGGGAGGGATGGGAGATG
GTGAATAGGTGAGAATGTAGGTAAACCTTCCAAGGTGCCAGGTTGATTTTTTTTAAATAGTAGCTTATATTTCCCTCAGG
CCAAGCCTTGGAGATTTTCTCTCAGCTGGTGGTTCTTGTTTTTTTTTTTTTTTTAAATTATCATTTTCTACTGCAGAGA
AAAAGCAGACCATTTCCAAGGGCAGCCTAAGGGAGCTGGAGGCAGAGAGTATCAGAGAGTGTTCAGTGCTGATAACC
AATTTTATGGATCAGTCATTTTAATTAATAAGGAGAATGGGGGAAAGTGATGCAAACAATGTAAGTCTGGTTGGCATT
TCCTTGAATGTTGAATACCTCTTACTTTTCAAAGGGTAAGGAATTTGGTTAGTGACTGGAAACAGGCAGAATTGGGGTT
GCGCTAAACTCAACCAGAGGTCACAGAGTACTGTTGGCAAAGGTTGGCCTCTTTTTCTTGTCTGCACGTGGCTCGTATTT
AATATACTACTGCAGAATAACTTTGATCTCTCTGCCTTTAGACAGAAGTCACCACCCTATCCCCCTAAAGCTATTGGC
TAGCATTTCTTTAAACAAGCAGGCTGCACAGAGCTCCTCATGTGACTCCAGCAGGGGAGGAAGGGAGGAAGTTGCATGG
GTTGGACACCCAGAGCTAAGAAGTAGAGAGATGTAGGTAGAAGGGCCAGCCAATTGGCAGCAGTAGGCTGCAACAGCCA
CACACTTGCCCCGGGAAGTGGGAATAGGAGGGATGCTAGAGCTTGGTTTCTAACATGGCAAAGATCTATGAGAGCGAG
AGAGAATAGTAATGGAGTAAACCAAAGGAAGAAATAAGATGCCTCCAGAGGGATATGGCAGCTTTAAACATGGCCTCA
AAATCCTTTGACACTCCTTCCATCAAGAAATGGGGTCTATGTCTCTTCCCCCTTGATCCTCTTGTGACTGCTTGACCAA
TGGGATGGGGTAGAGGTGTGAGTTTCTGGTGACACCTTCAGAAATGGGTGCGTTCTACTTTGTGTATCCTGAGAGGCT
TGCTCCCAAGAACCCAGCCAACATGTTGTGAAGAAGCCCATAGAAAGGCCGTATGGAGAGGAACTGCAGTCAGCCATG
TGTGAGAGTCATCTTTGAAGCAGATCCTTCAGCCTGTCAAGCAATTCAGCCGACATTACATGGCACAGAGATGAGCTG
TGCTTCTGAGCCTTCCCCAAATTGTAGATTTGTGAGAAAAATGAAATCACTGTTGTTACTTTTCACTACTAAGTTTTAA
GGTAGCTTGTATATGTGAACGATTGTAACAGGGTGCATTTTAGGTGACTAGGAAGAGAAATACCAGGTTTCATGCAA

Fig. 9.50

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TATTGAGGTTGCTTTGTTATTCAAGTATATAAAATATAAATCATATGATTAAAATTTTTTAATCATTTAGAGTTTTTTTT
ATTCTCTCCTTCGAAAAATGCACATATGCCAACCTTTTGAATACAGTTTCAAGAGTCTGAATACCTTCTACATCCATCC
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AAGCTTAGGGAAGGAAGAGGGAGAAGACACTGGACTTTCTATTGTTAGACTTGTATTTTATTCCAATCCTTTCATAGAT
TTGCTCTTTGATTGGAGATAAAGTGTAACCTCTCTATATCATATTCTATATGATATTCTATATCATAAATTT
CTTGGATGGCAAAATGTGGATTAAAATTTCTGTCCTTTCTTTCCTCAAAGGAATGTTGAAATTAAAAAAGAAAAAGA
AGTGAATATGCTTAGAAAAGCCATAACTTGCCCATAGAGGTACACAATGAATATTTGTTGAATTTTGTAGCTGTCGTTA
TTGATAATGGATTACTTAATTAGGTCGCTGTGGTGTATGATATAGATACTTGTTTTCATCCATGGTTCCTGGCTTCTA
GCTCCCATAAATCCTTGFTATAATGTTGGGGCACTTTAGGCCTCAGGAAACGGAATCTTTCCTCTAACCTTCTCCTGTC
CTCCTTTCACTTGCCCAAGGCAGGACTCTAATCTGATTGTGCGTCAAATAACCCTCATTCAGATCCTGTCCTATGCAG
GCACATGGATGAAGCTGGAAGCCATCATCCTCAGCAAACAAACAGAAACAGAAACCAAGCACCGCATGTTCTCACT
CACAAGTGGGAGTTGAACACTGAGAACACATGGACACAGAGAGGGGAACAATATACACCAGTGAGGGGAGGGAACTTAG
AGGATGGGTCAATAGGTGCAGCAAACCACCATGGCACATGTATACCTGTGTAACAAACCTGCACGTTCTGCATATTTAT
CCCATTTTTTTTTTAGAAAAATAAAGAAAAAACCCCAAAAAACCAAAATACCCTCATTCAGAAAGAGTCTGCTCTA
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CTTTTTGTCCAATCACATTTCTACAGGGTTGTCAATCATGTTTATGTAATGAAGCCTCCATAACAACCAAGAGGATTG
GGTTTGGGGAGCTTCCAGATAGCTGAACACGTGAAGGTTCTTGGAGGGTGGTGCATCTACGGAGGACGCAGAAGCTCAT
GCATCTTCCCTCATACCTCACCTACACATCTGTATCCTTTGTAATATACTTTATAATAAACTGGTAAGGGTAAAAGTG
TTCCCTGAGTTCTGTGAGCTGCTTCCAATTCCAGTTAATCAAACCCAAAGAAGGGGTGATGGAAACCCCAACTTGAAG
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CCTAGAACAAGGTCCCCAGGGAAAAAGCACCAAGGGGTATAGGAGTGGAAAGAGGGGGTCAAGTACAGGCATCTCTGCC
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CGCAAAGATTGGTATCAGTTTACACCAAAACAGAACACACTGCTGGTGAAGCACAGGGAGTTGGAAACAATATACACAAA
AGTCTCACAAAATAGAAAAGAACATGGATACTGGCTCTAACATTGACGTCAATTGGGATACAAAGTCTATTTACATTGTT
TCTTATGAGGTCAGCAAAACATTTCAAATAGCAACTCCTTGGCTGCCTGCAGTGGGTCTGCAAATGTGAAATAAAACAT
GAACTGCAGACCTGACTTATGAGCAGGACGTTCCAAGTTTCCTTCTTTGCTTTTTCTGTAACAGATCACCTTGTCTTG
TACTGCATTGGTAGATTTTTATCATCTCAGATGAAAGAATCCTAGTGTGGCCAGGGCAAATCCAAAACACAGGTGAT
GAGAGGCAACACCCAATGTTAAACCTGATCCACTGTGGTTATTTTTCTTTCCCATTTGGGAAACTATAAGGATGCAAT
GGGCAGGAGAGTAAAGCAGCAATCACGTCTGCATGGACTGAAGGCAATTTAGTTTCTATCAGACATGGTGACAGTGAT
CAATGCATCACAAAATCACAAACACAGATGCCAAGCACAACTGTGTACAGATCCCAGATCAAAGATATCTACCATA
TGTTTAGCTTTTACATCAATGTACATCAAGTCAGTTTGAAGTAGAAGCCTTTAGATATACAGTTCATTACTTTCT
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ATAAATGCATTGCTTGTGATGGATCCAGTGAAAGTTTATGCGCAGGGCTGCCCTGCCCAATGCTGCACTGGCTTTCTG
CCCCATCAGTGCTGGAGCTGGCAGAACCCCCCTGCAGAGGAACGGAAGCAAGTGACCTAGAGGGGCTGCAAAGAGTCT
TAGGAGGGCAATTACAACACGTGTTTCTTCTTTTAGGAGTATCTAATTGGGTTCAATGTGAAGGCAGCTCACTGTCAG
GCACATAGAAGAGACTCTGAGTTAGAGAGGTTGAAATGTGGTCACAAAATATCCATGAATGTATTAAATCCATAAGGAT
TTATTAGGGGCTCTTAAGCATTTGATTCCCTTTAAGGAAATCACTGCTAATAAAGAGGCACACAGCTTGAGAAGCTGAA
GCAGAGTGAAAAAATTGATTTATAGCTCATAGGTAGAGAAAGCAAAATTCCTAATTAGTCATTTTTTTTTTAAAGCACG
TACTAAAGGGAAAATTACTATATATTTTTTTTCCAAAGTTTCCTTGGTCTGGCTGTTGTTTGCCATTTTAAATTACAGATT
TATTATTTATTTTGTATAATATATACTCAGGAATTATTATTATTTAATTTTTCTCTTTTTTTTTTGCATTGGGCTTGGT
GCTGGGGAAACAGAGACAAATTATGCAGAGTCTTTCTTGTGTTGGGAATTCACACTCTGGCAGCCTGGTGTAGCCATGTG
CACAAACAACTACTGTGCAATTTGATGAGAGAGACAGAAGGCAGGGCAGTGGCACAGGTGGAAGTGCCAGCTAGGGTA
CAAGGAATTGAGGGAAGGCTCTTGAGAGAGAAAGAGATCTGAGAGCTGTATCTTCTTAAAGTTGGTGTATTCCAGGT
AGATGAAAATAGGATTGGGAGACAGAGGTGGACATTCCTGAGAAGGGGGAGTAGTAAGTACAAAACAGAGGTGGCAATAG
CATTATGGACTTGGGGACCGTTAGTAGGTTGATAGGATTATAGCACATAAATATTAGACCAGAAAGACTGAGAAATGAT
GCTGGAGAAGAAAGTAGGACACAGGTGATGGAATGACTTGCAGGCTGCACCAGAGGCACTTGGATTTTATGAGAGCCTT
GGAAGGTTTCTGTGGCAAGTGTGCGCAGTGCCCTCTCATATTCCTTGGCAGTCACTGTTACTGTACAAGACAACAGC
TACAAGTCTCAGAGACTACAAGTTTGCAAGACTGTAATGAGGCAGGCCCAAAGCACCAAGTAAACATCCTTGAGAGC
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TTCTTAGCTCAGGCTCTACTTCTGGTGGAAATCCAATCTGGGACAGGTTTTTGGCTTGTGGAGGAACAGTGGCAATGAAG
GCGGAGTCTGGATGATTTTTCATGAGAATGCAAGAGAGAGTGGAAAGACTCTGGTTCAATTTAGAAAGCACAGTTTCGTTGC
AATTATGGCCAAGCTGTATGTATTCTGGGAGACAGGGAAGGACTTTTTCGATGATTCAAGGTTTAGGTTTGTGACTGGT
GGGATCCATTAAGCAAAATAAGGAATACAAGAAGAGGAATAGGATTTTTTGGTGAATGTGTGGGATTGGAATTTGATTA
TGCTGAATTTGAGGTATTTCCATGGACAGTGTAGGCATTGGGGAAGAGGACTAGTTGCGACACCCCAAGTTTGGAAAGCTG
AGCATCAGCCCTGAGTAGTAGTAGAAGAAGTGGGTGTGGCTGTGGTCACTCAAGCTGAGTGTGGGGATGGGGAGTAGG
AAGAACCAACAGAAACAACTTCTGGGAGTACCAACCTTTAACAATCAAATAGATGAGGAGGAACCTTGCTACAGTGAA

Fig. 9.51

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GGAGTGATCTGTAGCACCAGGCAAGGGCCAGGGGAGTGATGTCGTAGAAGCCAAGAAAGGAAATCATGTCAGGAGGAGA
GAGGAAGCAAACAGAGTCAAAGACTGCAGAGAGGGCTTGGCAAGGTTCAGACTTCAGGCTCCCATTGGATTTGGAAGTTGA
GTGGTCATTGTTTATTTAAGAAGTTCATACTAATTTGGATCAATTAGAAAAGTAGTAGAGATTACCTCCACTTAAGACA
TTTAGTTTTATTACTTTGAAAAC TAGGCTAACCAATAATTGCCTAGGACAGGGATTGCTTTAATGAATAGGTAAGAATAA
TTTCTAATTATAGGGTGGCATGATTTGCGTTACCATACTGAAAAATGCTGCCCCCTCTATTTGTATGATTTGTATGAAGT
TCAGTACTTTGGAAGCTTCAAGTTTGTTCACCTAATTAGGTAAGGGTTTTTCTTTTAAGCTAGTTGCTGTTTTTTAGTT
TCAAGTTGGTCCCTATTTCTGTTTGAATTATCACAGGTTTCAATTTTATGTTGCCAAAAGATAAGATTTAATACCATAG
CTGATGCCCCCTTTGAATTTTGAATAACAATTCTATTTTCTTTTATACTATTTAGTTATTTGTGGTTAAGAATCTGGTCTG
TAGAGTTAGACTGCTTAGCTATTACAGTAGCTTCAACACTTACTCTTTAAGATCCCTGGGTAGGTACTGTGCTTCTCTG
TGCCTTGATTTCCCTCATCTGTAAATGAGAGAAATACAACCTTTTTTTTCAGAAGTTAATTCTTAGTACCAAATGAGCTAA
TTCATATAATGTATTTAAAGGCATTTTAAATGGCACAGAGTAAATGATCAGCACATTTTAGCGTTAGAAATATTTGTTA
GTATTTTTTCTATTTATAATTTGTTACTATTAATAAAACTGGGAGGCAGAATAGTATAAAGTGATAATGATTGAACATA
TAAACAGATTAGGGTTCTGCTCTCAAATTTTCCGCCA ACTCTCTGAACTGGGACAAGTCCATTCACCTCTCTGAGTTC
ACATTTCTTCACCTATAAAAGAGGGAGGTTAGAATAGAACATCTTTCAAGCCAGCCTTAGTTTTAATATTTCTGGGAAGCA
CACCTTTTGTCTTGGAAGTAGCTGATTTACACAGTAGTCTTAGCTGTAGTGTGTTTTCCCTAAGGGAAAGATAAATGGG
CTAAAAGGAGAAATGGAAGGCATTTCCATTAGGCAATTGCTTTCTGGAAAGTACAATAATGTTTTGGTTGGTTTATTT
TCATAGAATTCATTTCTTGGGCTAAAACATAAATCAGTTACTATATTCAGAGGGCTTATCATTTCTTTTTTGAGTATAC
TTATGGTTGGAAAAATTTCCAAGTGTTTTTTCATTTTTTTTGCCCTGTTTTTTCATACCTTTTACAGAAAAATAATTGTTTT
TAGTGCATTCATTTCAATTTGCTAATGTTATAATCTGTACAAAACAGCCACCTAATTATTTCTTTTAGATGTTAAACATG
ACACATGTAAATAAAATGATAATTATAGAATGTGGTGTTTTTCTTGATCTTACATTTTTTTAGATCTGAAAATTGGTCCC
TGTTCTTATTCTGCATGTACTCCAGTGAACTTTCCCTTGATGAGTTATTTTTTCATGCGCACACATGGGGGAGCTTTTG
AGACACTAGTTTGAATGTTACACTTTGAAGACTTTCTCAACAACCTTGACCCTAAGATGATGGACTGGAACCTTTCATTG
AAGGAAGTGTAGCAGGGTGCAACTGGCTGTAGAACTGACTTCCCTAGCGCTCACTGTCTCATATGCAGCCCTAGAACCA
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CGGTGAAATGGGGGAAGTCAGCCAGAACTCTGATTTTGACAGCAACATTATTTAAAGGAGTTTCTGAAATGATCAAAC
ATGTTCTTCCATAGAAGAATAAAAGTCTAAAAGCACATGAGAATATTTTTTAAAAAATCACCTCTGGAGGAAGCAGAAAC
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TTGTTTTTTTCATATGTAGACAGCCAAGTTTCATCCTTCACAGTTTGTGTTCAACTCCTTGCTGTTTTTTTTCTCTAATCT
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TTGTACATTTTCTATTGCCCATTTGGATGTTTCTACCTGGATATGGAAGTCAAATGGAATGTCAGCTACCAAATTTACCT
TCCATCCTCCACTTGCTTCCCTCGGCTTCTTTCTTTACCCACCTCCAGCTCACCAGTTCTTCTGACTTTCCCGTATC
TGTTCACTAAACTACCATTTTCTGAATCACCTGGCCTTGGAATTTGGTGTCTCTTTGACATTAACCTGTCTCATGTA
CACTACTGTATCAGGGGTTCCCAAGGCCACCCTCAGACTTGCTAAAAGGATGCATGGGACTCAGAAAAGTTGTTATAGT
CACAATTATGTTTTACTTAGTGAAAGAATACAGACTAAAATCTGAAAAGCAAAGATATGTGGGGAAAAGTCCAGGAGA
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GCATAAACTATCTGATCAAACCTGGTACCACATGCTCCAAGGCCTGAGGCATACAACACTCTTACCAGGCAGAAATATACC
TGTGGCTCAAGGCTTACTCTCAGGAGCTGGCCTAAGTTCAAGTCTTGAAGAGAGGTTTTTCTTGGGCATGTGCAGGGCTT
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CAGAGCACATACTGTGGTGTAGATATGTGTTTGAAGTTCTGCCTTTGAGACGATTAAATGAGATGTGATTAACTACTG
CTGGACACATAAATACTAAGTGATGTCGATTTCTCGTTGTTGTGATTTTTTATTATCTCTGCCCTTTTCTATCCAATATC
ACTTCTTATGTTTAGCCTAAGACCTTATTGTGTTTACCTTCAAGGTGCACTAGCTTCCCATATCAGTGAACCTTAGGATG
TTACGCTGCGCTAACAACAATCTCCAATTTGGTATCACTTGCTAAAACAAGAGAGTTATTTCTTGATGCTGCCGTTTG
TCTATTATGAGTTGACTGCAGCTCTGCTCCAAGTATAATCATTCTGGGACCCCGACTGAAGGAACAGCCCTGATCGGG
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CCTTGGAATGTTTGAGCAGTAATATAATCTATCAACTTCCCTAGCTAGCCTCTGTGTTAAATGATCATTAGAGTATCTAA
AGAGGCTGGAAGCAGAGAGGCCAGCTAGGATACACTAATCTATCCTGCTGACCATCACTAAGTTACATCCTCAAAGTT
CAGTGTTGACAAACCATTTCTGTTCTTATAACTGCCGTATCATGGTCTTCTCTTGATTAGTCACGTAAGCTCATTTTT
AAGCTCAATCACCCATGACTTTCCTATACGCAAGTGAAACCATAAAATTAACCAGCATCGGGCACATTTTATCTTAGCA
CCTTTTATACCTTTGCTCATACTTTCACTGAGATTGAGGGCTCTTTTTTTGAAGATAGTTACCATTTTATTACCTTTGA
ATTCTCCAGAGCCCAGGTACCTTACCTGTGGTCATGGCTGAAACATGTAATCAATTAATGTAAGAAGAAGTATTGTCCC
AGTGTTCAAAGAAATCCAGGTGCTATTCAATGACAGTGAGACGTCAGGCCAAGGGGGTGAAGGGAGGCTGAAACCCAG

Fig. 9.52

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TCCATGCTTTGCTCACAAGCCAAGTCCTGAGATGGGATGAGGAGAAAGAGGTGTCTTTTTCTAATCTCATTAAAGCACT
GAGTAGTGTGGTGGGGGCTTTTCTTCTGGGGTTTTGTTTACTAAAAGACTTCCTACAAAGAACCCTGTAGGCCCCACAAA
GATCATATGCATGGACATTATTGTAGGGCAGCAGGAGAAAAATGCTATTTTGGTTCTGCTTTCTAGAAATTTTCAAGTG
CTGGGCTACCAAGTCAACTAGCTCCTCTGCATCCTTTAGATGTCTGTGGCTGAGGACAGCTTCATGAGATTGGGTCTC
AGAGCTGCTTTGCACTTCCCAAGAATAGACCTGTGGACCATGTCCTTTTTGTCCACCCAAGTTTTATTATTTTTTGGGA
CAGCACCTCTTTTACCAGAGAAAGTAACTCTTGCGGCTAAAAATATACCGGAAATAAGAATGAAGAAAAGTAACTGGAT
CAGCTATACTTGGTAAAAATACCTAAAGCTCTGTTTCATGAAAGTGTTCCTAAAAATAAAAACTAGTCCCTGGCAATGC
AGCAAATAGCCAAAACGACTTCTTTGGTTGATTGGTTTTTAATGTTTTTTTTTTTTTTCTTTTCTGTTTACAGTTTTAA
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AGCAGAAAGGCATGCAATAGGGCTTTGATTGTCCAAGGCAGGGTAACCATGTATGTAGGACCTTTATTGCTGCCTGGGA
GGTGCTTCAGTGTGGGGAGTAAATGCTCAGTCAAATTTGGCATAACAGCTTTAAAGCAAGTGGAACCACTGTGCATGTT
GTCAGCAACATACTGCTTTCTTTCTTTCTTTCTTTCTCTCTCTCTCTCTCTTTCTTTCTTTCTTTCTTTCTTTTAGTTTT
TGAGTCGGAGTGTCTGCTCTGTGCGCCAGGCTGGAGTGCAGTGGTGCTCTCTAGGCTCACTGCAATCTCTGCCTTCCGGG
TTCAAGTGATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGATTACAGATGGGCACCACCATGCCCTGGCTAATTTTTGT
ATTTTAGTAGAGATGGGGTTTACCATGTTGGTCAGGCTGGTCTTGAACCTCCTGACATCGTGATCCACCCACCTTGGC
CTCCCAAAGTGCTGGGATTACAGGTATGAGCCACCATGCCCTGGCCCTTTTTTTTTCTTTTAAAAGTATTTTACTTATGTT
TCTATTTTGTGTCAAGGGGGCTGGTTCCTAGATTACTGAGGGAATAAGCCAAACAGAGCAAAATTTGTGTCACTCATAT
TAAGGTATTAATTTACGCACTAATGAGATGTTGAAAAACATGGCAAAGGTGATTAGAGGATATTGAACGAAGGTAAATT
AAGTAGGTAACTTGGATCCTTATGTTTCGGAATCCTGGTATTAGGAGTAGAGTATGGAGCTAATTGCACTGTGGTTGGA
AAGGGTTTTAAATTATAGAGAAGATGAAGTTTGATCAACCCAACATGTGTCCCTTTACTAGTGGGAAGACACAAGGGC
ATTCATGTCACCATGAAATATTCTTATTAATTTATTGAAATACCAGTGATTTTGTGGCATTTTGGGATTACTATCATTA
CCATAACAGAACTGTTCTTGCTTCATATTTATGCGCTCCAAGATTGCTCTAAAATTCCTTCATATCCAACCTGGGCAATT
ATGATTCACCCATTGGACAAAACAATATAGTTCTATTCTAGCACTCAAAGGATGTAGTCATCACTTCCAAGAGTGAGCA
GACTTTCTGGAGCCAGGTCAGAAAGAGACATTAATGTTGTCTTCATTTAAATACTATGTGTAGAAAAGACCTATGTATT
TCCCATTGAGAAGATGGATCATCTTTTTCAGGAATTGTTTTTTCATCTGTAGTCTCATCAGATCCTGTTAATATATTTTA
CTTTGCCTTTGCTTTTTTTTTCCATTTGAATATATCTTTTTCTCCTTTCAGCATGATGAAACGTCATTTAATTATAAATT
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TACCTCTGGGATCATTCTTTGCAGTTCTTGGCTCTTGGTTTCTGTAGGAGTTGGCAGGTAGATGCCAAATTTTCTGTCT
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TGGCTAATAACAATATTTGGTCAGAAATTTGAAGCTGATAGCTTTCTCTCAATGTTCTAAATAGATTCTATTCTTTTAA
AGTGGTTAAAGTAGAACAATCCTTCAATCCCTTGATCTGATTTATGGCAGTAGTAATAAAGAGGATTGTTACCCCAAAT
GGGCCAGTTGTTTATGCTTGTATTATTCATTTATTTCTGGAATGTTTGAGTGTCCACGATATGCTGACTCTGTGCTAGATA
CTGGGGGTTTCAGTAGTGAGGTCCAAGTCAGATATAGTCTTTGTCTCATAGTGTGTAAATAGTGAAATCTAATGGGGA
ATACAGAAATAGGCAGTGACAATAAAGTATTGTGAGGAATGCTATAGGCGAGAAGCACAGTGAGCTCACAGGAGGCA
CATGGACAATATTTGGTTGCCTGTGTTTACCAGGAAGTGTGCTAAAAGCTGAGTTTACAGTTAACTAACAAAATAGACA
CTGTCTTGCCATCGTGAAACTTACTATTGGTGAAAGTCAGTGTGACTGAATAATTACATACATGCATACGTACGATG
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GAGCCTATTCAACCACAGGCAAGTGTCTTGACTTTACCTGGAGGGGAATGGAAGAGCTACTGAGTGTTTTTAAACATGAA
GGACATATGAGCAGATTGATATCTTTGGAAATTCGCTGTGGCTTCTGTAGTAGATTGAAGTGTGGGATGCGTTAGGAGG
CCTGGAGACTAGGGAGTAGGAGGCTGTTACAGGAACCCCGGCTAAACAGTTCACTATAGAGTAATGCTGGGAGTGTGAG
CTCTGGGGCCACAATGTTTGGGCTTAGTCTTACCTGTGTCACTCAATAGCTGGGAAGCTCTGAGCCATTACTTTTCTTC
TCTGGCCTTTGTTTTTTTTGCTTCTGTTAAATAAGGATGATGATAACAATGATGCTTATCTCATAGGGTTGCTGTGGGTTT
ATCACCATTTAGTGCTCAGAGCAGTTTGTGGCCCATTTCTGGGACTCTGATATTCATAACTATAAGATAATAAATTTGTA
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CAGTTGTAAGCTGTAAAAATCTGGGAATCAGGAGATATATATACGTATGTTTAGATGGAGTCTTGCTTTGTCAACAGGC
TGGAGTGCAGTGGCGCGATCTCAGCTCACTGCAACCTCCGCTCCAGGTTCAAGTGATTCTCCTGCCTCAGCCTCCCA
AGTAGCTGGGACTACAGGCGCGTACCACCACACCCAGCTAATTGTTGTATTTTTTAGTAGAGATGAGGTTTACCATGTT
GGCCAGGATGGTCTCAATCTCTTGACCTCATGATCTGCCACCTTAGCCTCCCAAGTGCTGGGATTACAGGCATGAGC
CACCATGCCCAGCTGGTAGTATATTATAAAGAAATGATATTAGATACACTGTATTTTGTGTAGCCTGTGCCCTCCAAA
AATGCTCATGGTTGAGGGGCCAGTACAGGTAGGAAGTGGCTTTCACTGATAAGCCCTGTGCCTAGGGAAGAGACTCCTT
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TGGCACAATTAAGTACTCAATAAATGATATCTGTTACTAAGTGTGTATGTGAAATTAAGTGAACCTTTCTATGCCCA
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AGCCAGCTATGCAGAAGCACAGGGCTGTAGCTGGGGAGAAAGAGACAGATATGTCCCTGTCTCATAGGGTAAACAAAA
ATTTACCCTGATAATGATTTAAAGCTGTGATGTGTTATAAAAATGAATTCTAGGAAATGTAGACAATGAGGAATATTTG
AGAAAGTTTAGAAGTAGAAGAACAAGACTCTGATTAAAGAAGCTGATTCCAGAAAGATTGGAAAGGTTTCAAGAAGGA
GGGCAAGGTTGGCCATGTTGCCACCTTCAGAGCTAGAACCAGATTACCTAGCTAAGTGTCTATGACTTAAGAAATG

Fig. 9.53

Fig. 9.54

Fig. 9.55

ATCTTCTGACATTTCAAAAGGAGCTGGAAATCAAGATGTGAAATCAGATTATTTAATATTGACAATTAATTCAAATGTT
TGTGCTAACATTGCAAGACTTCTCCTCATGTTGTTGTCATGCCAAGATGGAGAAAATATGTTGGGCACGTTAGGTATTTA
TGTGTCAGTTTGTGACCTCTGCTCTGGTATATAAGTTTGGCCTTAGAAATATGTTCTGTATTATTTTAAAGTTGAGTG
GGAAGCTTCTCTACTTTTAATGAGTTTAAAGGAGAGATGTCATCTACTTATCCCTCTATCATTGGAGGATAGGCTAG
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ACACTTACATTTCTGCTTCTTGGGATAGAGGTAATCTGTATATTTTGGTTTTTAAACAGATATTTAATCTTTTAGGCA
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CACTCTGCCAAGTTGAAAACAAATCTAGGTAAATTTAAATTGGGCTGGCATTTCACAGAAGTGGTGCAAGCATGGTTCA
TAAAGTTTATTTTTTTGCTTGAGAAAAAATGACATTTTACTGATTCTGTGTAAGAAAACCTTTAAGTTTTTATCAAAG
ACTATTAAGATCAGAATTCAATATTTTATGATAGTATTTAATATTTTACAATGTATTAATTTACAATTAATAAGTTAT
TTAATACAAATAACTTAATATTTTACAATAGAGTTGATTATTTTTCTAATGCTTTCCTAAAGAAATTCTGTAAAGATCC
ACATAATTGCTAGACCAGTGATAAAGCATGTAATTTCCCTTATCTTCAATTTTTCTTACTGTCTACCTCTAAAGTAG
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AGGATCCCTCAGTTTGGCTGGTCTTTCTGAAAAGACAAGTGTGAGCAGCCTGAGGAGAACCTGGAAGTCAAGGTATTCTC
TGTTTTCAAAGTCAAGGAAGGCTGGGTGAATTCCTGGTAGAGCTGGTAGAACTAGCTTTCTATTATGAATATTGAGAAGT
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GTAACCTCCACCTACTTGGGCCAGCCCATGTAGTTCAAGGTAAAGATCTTGGATTTTATTCCACATAAGGTAAAGATCT
TGAGCTACAAGATCTTGAAGTACTGAAGCTCTTGAATTACACGGGCTGGCCTCTGCCTTTCTCTCCAACATCATCTCCA
GTGACACTGGCCTTCTTTCTCTTCTTCAAATAGACTAAGCTCTTTCTCACTTCAAGCTCTTTGTTGAACCTGCTAGTCC
CTGTTTCTATATGGCTCTTTCTCCAGATCATTTGGCTTCTTTTCATTTGGGTCTCAAGTCAAATATGACCTTCTTAGAGA
GGTCTCCTCTGGTGATGCCACTTCTCCTATCACAGTATCCCATCACTCTGTTTTATTTTCTTCATAGCCCTTATTTGAA
CTTATCTTACTTATTATTTGCCTCTCATTAAATGTAAGCCTTGTGAAAGCAGTACCTTCTCTGTTGTACTTTATTGCT
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CCCACATTCAGTGGCTTTACTTGTAAATCTGTTTGAAGTCTTGAATTTTATTATATGGAAGGGATGGAAAG
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AAGTACGGCCTTAAACAGAAAAAGCAAAATATATACTACTTGAAAGTCAGACTGTATGATAATCTAATGGTAGATA
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CTCAAAGTCTGACCTTAGGTGATCCACCCACCTCAGTCTCCCAAAGTGTGGGATTACAGGCGTGAGCCACTGTGCCT
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TTAACTAATGCAGTAAATAATCCATTGAGATTTAATCCTCATCAATCATCCTTATTGTTACCATGTGGCAGAAAAA
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CAAAAGATGACTCATCTGACCTTCTACTCATCTTTTTTAAGACATGAAACAAAAGAAAGCAAATGGGTCTCTGTGGCCAA
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TCTCTCCTCACTATTCCCAGCACTGATTACAAGCTTCAAGCAGAAAGCAGAGTTTTTAAATCTTGTGGAATTAATAAATA
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GGCTGTGACATGGTCTTGGCTCACTATTATTTTTTTTACAAGTTTTAGGATCTTGGCGTTTTATTGTCTTTATCAATTAC
TGTGGACTAAGATGTGCGACTGTGATGTGGATTAAACAACAACAATTTATTTCTTGCTCATGCTGTCTGTCCATTGTGG
GTCAGCTGAGTGCTCTGTTCAATCTTGTCACTCAGGCATGGGTTGATGGAGCATCCACCATCTCAAACGTTGTTAATTA
CCATGCTTGGGAAGAAAGGAACTCTAAAGGATGTTACCCAGGTGGTTAAATAAGCTCATGTAGAAATGGAATGTGAC
ACTTCAACTCAAAGTCAATGACAGAACTCAGTGTGCCTGGCCCCACCAAACCATCCACAAGGGAGGGCAGAAAGTC
TAATTGTGAACAGAACTAATGATATTTACTTAGTATTTGTCTGCTTGATTTTTTGGCACAGACTGATTTATTAGAGTGA
GAAGGTAAAGCTGATTATCAAAGAATGTCTTATTTTCTCATGTAGTTACCATTGATTTATTGAATCATGATTTTCAATCT
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GGGAACAGATCAGACATAGGGAAGGTGGGTCTCATGTGCTTGATCATCTACTATGTGCTTGATCACTATGCTAGATACT
ACATTGTATGGCTTTGCTGAATTATTGTTAAGTCTTCACTCACTTAGCTGAGCTCAGAAATGAATACATCTCAATGCAT
GCAGAGAAGGTTCTTATTGAGGCTTACTCACTTTGTAGGGTGTGGTGGCGTTTGAGATTGTGTTTTCAAAGTGCTTAG
CACTGGCGATAGTGATGTTTCAATGATGTTGCAAGTTGACTAGGATTTGATAGTTTCAATTTCTAAATAATTTTTATT
CAACTAATAATTCAAGTAAACACTGCTCACTTGATGATTAGTCGTGGAGTGAAGACTGCGTTGTGAAGCTCACCATTC
TTCTCTAACAGTGGTCTCCTGAGTTGGGACACTTACATAGGATTTATTTCTGATGATAACAAGAGCACAGATGCCAACC
ATGCATGTGATAGTTTCTGGGAGAAGACTTGATGATTCAGCACTGTCCCTATAAAATGACAAAAGAACCCACATATA
GCATTAGGTATGGATGAAGGTACTTAAATTTAAGCTTAATTAGGTGTAAATCCTTAACTCCTATATTCTACTCTCTGGT
GCTTTGAAGTTGGCCTCTTCGGTCTCCAGCCACAGAAGGGATTTTCTTCGCTGACCACAGTTCCCCACGTTTTCCCTTC

Fig. 9.56

AGTAAATAGCAGATGTGTGTGGTGGTAAGCTGGTTTTCTGCCATTGCTGTGTGAAAAGGCAGATATTCTGAATGAGGAT
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CTCTTGATGATTTTCTCAGATTAATGCTGTCATCATGTTAATGGGGTAATGCTTCTCTGCAAAGATCTTGGAATTTA
GGGCTTCTGGTCAGTGTTTTACAGCTTTGTCTGAGAAAGAATGTAATATGAGTATGTTTTTGGTACATGACTTCAAAAA
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CTTACCCACTTTGTAGGGTATTGGCGGTGTTTCGAGATTGTGTTTTCAAAGTGCTTAGCACTGGCACAGTGTGCAACTT
GCAGCATAATTATGAAATACCTTAACAAAAGGTATTTACAAAAGCATGAGAGGATGAATGTGAGGAAGGAGCATGTCAT
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GNTGGATCAGACACCAATTTACCAAATGATTCCATAATATTAGTGCACAGATAATGCACAGATAGTGTGCAACAACAGC
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TGAAGAGGTTTTGAGCAAATATTTTAAGAAGTCTTTAGGGAACAAGCAATTTTCTTTTTTGTGCCCCCTTCTAAGAATA
AGACATAAATAGGAAGTTCATCCTTTTATTTGTTCTATTCAAATATTTATAGGGGAAAATCGGCTACTACTTCTTTAT
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ATTTATTTTATTGAGATGGGGTCTCACTCTGACACCTAGGCTGGAGTGCAGTGGTGCATCCCAGCTCACTGTAGCCTC
TACCTCCGGGGCTCAAGTGATCCTCCCACCTCAGCTTCTGAGTAGCTGGGATCACAGGCTTGTGCCACCATGCCCGGC
CAATTTTTTGTGTTTTTGGTGGAGATGGGGTTTTGTCATGTTGCCAGGCTGGTCTCGAACTCCTGAACTCTAGCAATC
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TGTTTAGGCTATGGAAGACCACATATATTCTTACTTAAGCACTTAGAATGGAAGCCACCTGAAGACAGAGGTTATATCT
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TCCAAGTAGAAAGACATAAAAGTTTAGTCTTTGAAATGCAGTTGGTCTGGCATAAAAAATAAAAATCAGGCTCTAGTCC
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GCATGGCATGGTTTTATTACATAACAGATGTTAAAGAGTAAGAGTGATTTTTTTTTTTTACAAAGACAACTCAGAAAATAAG
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TAGCAGAGGCTTATGAGAAGAGCTTGGTTCTGTGGAGCAGCAACATGATAAAGTTAGATTTTTTGGGGCCTCACCTAACC
TGTGGGACAGCTCTCGGGCCTCTGATTTCAATTATTCCTGCTTTTTTTTTTTTTTGGTATTGCTGCAACACATCTGTCTC
CTGGTACCCTCTCCTTGGCCCTTCTGTCTGCTTTCCACTGGGATTGAAATGCTTTTTCTTTCTCCTCCACCTCTGTTTCTAG
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AGTAGCTGTGTTGATGGACGTGAGTGTCCAGTTCATAGCTGCATGCCAGATGAATAGCAGTAATAGAGCTACAGAGGC
CCTGCCAGCACAGCCAGACATGATGCACTAACATGCCAGTTTCCTTGCTTCTGGTGTCTTCCAAGCCATGGGGCCCTGG
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AAATTCCTCTGAATTGAGAACCAGACTAGATGTAGGGATAAAGTGAAGTACAGTTGATCTTTGANCAGCACAAGTCTGG
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TCTTCTCCTGAGCCTACTCAATGCAAAGATGATGAGGATGAAGACCTTTTGAATGATATGGTTCCACTTAATATATAG
TAAATATATTTTCTTTTCTTATGATTTTCTTAATAACATTTTCTTTCTCTAGCTTACTTTATTGTAAGAATACAGTA

Fig. 9.57

Fig. 9.58

TGAAGCTCTTGAAAAAATTTAAGAGGAGGAAACAGAGTAGGGATCTTTTTGGGGTCACAGTCTAGGTGTTTTTAGAGC
CACTCAAACCTCGAATCAAGACCCTTTGATTACCATAAAAAATCTAAAATAAGATGTTTTGTTTGGCCCAAACAGTTTTA
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GTAACCTATGGGTTGTAGCCAAGTTGTTCCAACACAGGTGGAGTGAACCTCCTGTGGTATTTTGAAGTGATAATTGGCTAG
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CCTACTGATGAATGGTGCAAAATAGGAAAAAGGAAAAATAAAAAGTATTTGCTACCTTGAGAAATACACTCAGTCTCTATT
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AAAACAGCAATTGGCTTTTCACTAGTATTTGTTCATTGCTCTATCCTAATGCATTCAACATTTTCTATTGTGCTATGCT
CCACTGTATATCACATTTTACCTATTTGATTCCAAATTTACTTTGTGGTATTAGGTTTTAATTTAGTAATTTACATACA
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TTAACCTCTTTTTCTAGTGGTCCCAATTTTATTCTTAGCAAAAATAAAATTACAGATTCTGTTTTCAAAGAAAATTTG
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TGTCTTTTAAATATATGTTTCAGTTATAAAAGTAATATCAATTCCTTAAATAATTTTTGTAAATATACAGGCAAGCCGCAA
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TTTTTTCTACTAAGACTAATTAATACTGTGTGGAATACTTTTATATATAGACCATTTCCTACTTGAGTTTTTTTTTAAA
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AAATCCATGTTTCCTAATACTCTGCCTGGATTTTTTTTTCTGTGTTTGGAGCATCAGATGAGAAGTTGTGAGGAGAAAC
AATGGAAAAAGAAGAAGGAATCATAGAATCATTGCTGAGTGTTGCTCCCCAGGTTCTTGTTTGCTAACTGAAAGGAATT
TGCTGGCTGAAGGGAAGACATGTTGAACACTTTTATTACCCCTATATTCACCCCTGAATTCCTCTCCCCATGGCCAAGGGA
CAGTATCAAGACTTCTGTTGTTTCACTGAGACAAGTCAAATTAAGAAACACTTGAGTGTTTATTGTGTCCTTTGTAGGG
CTTATGGGCATATACCTCAGCAAAGTAGGCGGGCAATAATTTAGGTCATGCCCATTTAATTTCAAAGGCCAGCTTTAAT
CTTTTGGGCTGATTGTGATTACTCCAAAAACGGCAGAAAACAACCTGAGAAGGGAATACCAGGTCCAGGTTACAGTTCTA
TGGAGGTTTTGTGTAATCCTGAGTTTTTCGCTTTGCAATTTGGCACCATGATAGCTTTTGTATTTTCTTCTCTGTTCT
ACTTTTTCCCATCATGGCAGGTGAACTTCTCAGTGTGCAAGGAATTATATGTAGCTGACTCTTCCAGGGTATGAGAGGT
GGGCTGAAGCTATCGGGATAGCAGCTGGCTTCAGTAGTGCTTTTAGCTGTCTAGACTTGTGCTTCTCATTGATTTTGG
TGTCTGGGCATGTCCTTTACTTTCTAAATCATGTCTGCATTTAAATATTTTAGAATATTTTATCCAGCATTTTTTGTG
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ATGCAACCTATTCTCTACACAACAGTCAGGGTGATCTTTTAAAGTATACTTTAGATCATGTGAGTCCCTACTTAAGAC
CTTCCACAGCTTTTTCATTGCTCTTAGATTAAAGCAAAACCCTTTATGTTTTCTACAAATGTCTGCATGATCTGGCTCC
TAAGAATCTTCTTTCTTACCCACTCTTCTCCTGCTCCTTAACTTCCAGCCACACTGGCCCTCTCTGTTTTTTGAACTA
ATGGAGCTCATTCTGCTTGGGGCCTTTGCACTTGACATTCTTCTGCCCATGTCTGGACACACTGGCTCTTATGCAT
CATTCAGTTCACAGCTTAAGTGTCTCCTCAGAAAGGACCTACCCTGTCCACCCTACTGAAAATAGCAAATTTGCTATCA
CCCAATCATTCTAACCCTATCACCCTGTTTAAATTGTCTGCCTTAAATTGTTTTGTTTGTCTATTTGTTTACATGTTTACA
ATTCAGAATATAAGCTCTGTAGGAGCAGGAACTCTTCTCTCTAGTTTCACTTCAAGCCCATTAAGAACAAATGC
ATTGTTTGTGGAATGAATTAATGATTTAAATAATTAGCCTATATAGTGTCAATTTCAAAGTTGGCAGAATATCATCATT
GGAATTTTGTCTTAAAGGGGAACCTGGGTTATAGAAATGTGGGAAGAATACTTAATTAGCATTGAAAAACAATGCTTTAC
TAAGGAATCAAAATCATGAAATGTAAGAGCAAATAGGAGTGTTTATCCCTTTTATTCAGACTCTATTTTATAGATATA
AAAACCGAGACTCTGGGTGTGCTGTGAGTGGCTCAGACCCCAACAATTAGTGGCAGAGCTGAGGCCAGGGGATGCACATTC
TGATTCTGCATCATGGCTGTTGCTTAATATCTCACCATGAGTCATGGCTACTGCCCAGTAGATTGGGCACAACCATGA
ATATTAGACTGGCAATTTGCTAGAGATTTTGCAAAACCATGGGAAATGGATTGAACAAATTTTCTGTCATTTTTGTAA
GAGGCAGTAAATAATGGTAATTTGAGTATTAGAGAACAGCATCTGAATACTTTTTCTAAAATTCTACAAGGTGAACATA
GAAAATTGTAGCTTTCTTTCTGCCTATTGCTTTCTGAATGTCAACAGATTGCTTTCTTCAGCTTATGTTAGAATGTCA
CCAGCAGGACCCACATCTGACATGCTCTGGACTGTCAGAGCCACTCAGCACTAGGAGAACTTTCCAGTTGAATTTCTCT
TAAGAAGGTCAGTAGGAATAAGATAAACGAAAACTTATTCATCATTACCATAATCCTTGTGAAAGTGGAAAAGTTCAT
CCTGGCAAATTCAAAATCGATTACAACTCATGCATGTGTCATATGTATTTTTAAAGATTTTTTACAAACCCCAAATAAAA

Fig. 9.59

TAAGTAGAAGAAAACAGTAGTGATAAATTTAAAGTTCTTAACTGAATGAGTAGCTTTGAATTTTACTCAACGGATATAA
GCTTTGGTTTTTTTTTATAAAAAAGAAAACAGGTTTGTTAAGAGACTATGAACATATTTTAGAAATGCTTCAGTGATTTT
ATCATTTAAATAGAAATTTGGATTATTCTGGAAAACAGTTTGATAATATGTTTCAGGCATTTGAACATATTTACTCTCC
TGGCCCAGTAATTGCAGTTTTAGAAAATTTATCCTATTTTAGGAATCTAGCCTAGAGAAAAATCTGAAATTCAGTCATCT
TGGCATAAGGATGGTTGCAGAAATACTATTAATAATAAAAAGTTGAATAGTACCTAATATTCAAGTAGAGGTATAAATA
ATTTGTGGCAGGTTTCATTTGCTATAGTTTTATGCAACTATTTAACAAATATAAGTAAGATTATTTAAATGATGAAGGAA
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GTGCTGCACCCATTAACTCGTCATTTACATTAGGTATATCTCCTAATGTTTTCCTCTCCCCCTTCCCCCACCACCGA
CAGGCCCCAGTGTGTGATGTTCCCTTCTGTGTCCAAGTGTTCATTGTTCAATTCCCACCTATGAGTGAGAATATG
TGGTGTGTTGGTTTTTTTGTCTTGCGATAGTTTGCTGAGAATGATGGTTTCCAGCTTCATCCATGTCCCTAGAAAGGACA
TGAATCACCCTTTTTTATAATATAATGTTAATTTTTTAGAGAGTAAGACCTAAAACCTATATATACGTGTATGTAATAC
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CCTGTATTGCTTTTCTAATAAGACTGAGGTAGTTGGTGAAAATTTGAACAATTCCTTGCTATAAGATTTTATTGCATAGG
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TCACATAGTTGCCTATTTTGAAGCATAGACATAAAATTAATAAATTCATCCAGATAAGTACGAGGCTTTGCCAAAGT
GCCCCTTAAATTGCTCCCCCTTTTGAGAATGCACTATTTATTGTAGCATTTTGCTGAGCCCTTGGGAATTTCTCTTCTG
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GTGCTTCACTTTAAAGACTAAGATACTGACGCAGAGGTTATAAACCTCCCCCAGGTCACAAAATCATGCTGCTGGACT
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AATTACTACATTTTCAATCTGGTGCATAGTTCTGAGTTTTGTACATCCTTATGTGGCTCTACACTCTTTGAGGTTAATT
TTGGCCTTGGATGGTGCCCTTTTAAAGGCAGGGTAATAGCAACACAGTGTTTTTGGCTTGGGAAACGCTCTGTGTATGGG
CTTCTTCTCTTGGTTTTAAGTATTAACAAGGTAGTAAGTATGAAAGGTGCTGTGTTTGGAGTCTTTAAATGGACTTGGC
ATTTGGCTTGTCTACAATCTTTCAGGAATTTTAATCCTGATATTCAGATTTGAGTCACACACCTGGGGAGTGGTGACCA
TAGCTCTTCTTACCCCTCCAGCTCCACTTCTGGGTTAGCATGAACTGACCACTGTAACCTGCTCTCAGGCTCTACACA
AAGGTGCTTCAGAATCTGCCTGTGGCCACTTCTGTGAAAGGGCATGGTCACAGTCACCGTGGCATGGATAGAAAACCTGG
AGAGCAGCAGGAGAGCCAGGGATAGAGAGAGGAGGCTGGATGCCTGCCTGGGCTCTGTGCCATGCCTTGGGTAGGC

Fig. 9.60

CACTTGACTACTTTAAACCAAGTTGACTCTCCTGTAATGCAATAGGGGTTTAATGATCTCTGTGGCTCTAGAGTTTTGT
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TCAAGATATTGCTCTGACTCACTGGATGGCAAATCATAGTGAAAGGGAAGTCTGACTCGACTTCCATTTTGCATTACTT
TTGGCCAGCATGGCTCTTTATGGTGTACTTGTGTATAATAAAACAGCTNTCTAGAGAATACTCTTACATTTACTTACT
GAGAGATAACAGAAGTTCCGAATAAAAACTCATAGAAATGGAAGACTNGGAAATTATTTTTTTCTTTTTTCTATTACAT
ACAGAGGAACAAAGGCTGATATTTGTGTTTATGCCCCCTTCTGAGGACAATGTCCTTGAAAATCCATAFTATTATTATT
TGTCTTTATATGTAATTGGATTGTTTTAAACTCTTCTCTGAACCTTTGTAGCTTTTTTCATTATATCTCTTTTGCTGGAATA
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TTGTGGTTTTTTTTTTTTTTTTTTTTTAAATGATGTGACTGCTTTGTGAAGGGACATAATGAGTCTGTGTTCTCTTCTATT
CCTCTTCAACAACCTTTGTCAGGAAAGAAGGCTATTATGATTTCTTGGAGATGTGGGAGGATTGTGGCATCACCATGTCC
TAGTCATGGATGAAAGNAGAACTATTATACCAGGGTAACATCTGAGGCCTGAGATAAAATTTCCCATTAACAATCTCTTT
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CCGTGGGCCCCGATCCCTGTTTTGCTTTCTTTCAGAGAGCTCACTGGCAGCCTCCCTGATGCTTTGTGCCAGTTTTTAGG
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CAAGGCAGAGGCCTTTAGGGCAGGGATCTTCTGTGAGCTGAAATAAAAGGGTCTGGTTTGGAGGAGATTTGACTCTGCC
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CNAAAATATTGAGTGAACATTTAAGAGATACACAATTCGTAAGTTTTAATTTGCATTCTGTTCTGAGTAGTGTGATGA
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GTCCCTTAGTAGCCGTCTCATTAGCAGATCCACCTTCGAGGTATGGAAGTGCTTGTGTTCAAGGAACCTTTAGTTTACT
TAATAATGGCCCAAAGTGCAGGAGTGGGGATGCTGGCAATTTGAATATCCCTAAAGTGCTTCCCTTAAAGCTAAAAGGTG
AAAGTTTGAACCTAATAAGGAAAAAAACAAAATTTATATGCTGAGGTTGCCAAGGTTTATGCTAAAAATGAATCTTCTA
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TGGAACCTTGGAAGAGGCTGTCCTTACCTCCCTGTGCATTCTGGCAGCCCCCTGGCACTGCTCATAGATAGCCATGTGA
CTCAGGCTGGGACTCTGACTCTAGAGCGAATGGGTAAAGGTCAAGAGGCTGTTAGAACTTATTGTTGGTGTGAGTGGT
GGCGCCCATTTGGCTTTTCAAGTGGCGGCGGCAGCAGCAGTGTCTGGCTGTAATCTTGCTACAGGATAGTGGTCTGTGATT
CTTGCTCCTTGGCATCTTGCAGCGCCTGGGTTTTTGTCTCATTCTTCTGTCTTAGTGATGAAGACTTCTACTGAGTCTGAA
CCTGCATGTCTTCCAATGGAGTCTACTTTTTTCTCCTCCAGCTAGCCAAGGCCGGTTTTCTGTTATGGCAAGCAAGAAT
TCCCACAGCACCTGAGTGTCAACGGCTTTTCATCGTTTGTGCTGACCCACCTGTGCAGCTGATCGTCTGCCTTTTTTCTC
CGGCTACTCACCTTCCTTTATCTACCTATATAACTCCCTGCTTTGAAGGCTGAACTCCAGCATTTTTTAACTTTTCTCTGA
GTTATTGAAGAGACTTTACTGTTTACTATTCTCTGTACAGTACAACATCTTATAATTGCCCATTATACAGATTTTTTTT
TACATTGTATGTCTTTTTTTTTTATTATTTGTTTCTGTTTGCCTACTAGCCCATGAGGAATGTTTTTTTGGTAGGAAC
TCTATTTTTCATCTTTATAGTTCCAGATCCAGCTAAGATTACACTCTGGTTGCGTGACTGAACAAATCTGTTTGCAGGGA
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ACTAAGGGAGGAATAAAATGGTGTCTGAAATGTGTGAGGGCCGTCTGATACCGAGTAGTGTACATTTATTTTCCACTTA
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AGATGCATATAGTCATGTGTATACCACCAGAATTAATGTATAGAACAGTTTGATCACCTCCCTCAAAGCTCTCCCTC
CTTCTTATAGCCAGTGTCTCCCCACATCTTCAGCTCCTGGAAACAACCTGATCTATGAATTGATATGTGTGTNTGTGTGT
GTGTGTGTGTGTATATATATATATATATATATATATATATATATATAGTTTGTTTGTTTGTTTTTTGGAGACAAAGTC
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TATTTTTCAGTAGAGACAGGGTTCAACATGTCNGCCAGGCTGGTCTCAAACCTCCTGACCTCAGGTGATCCACCGGCCTTG
GCATCCCAAAGTGCTGAGATTACAAGCATAAGCCACCACATCCAGCCTATCAATTTATATTTTAAAGCAGTTATCATAGT
TCTACAACCTTTCCAGAATAATGTTTTTATACACCAGGAAGATAAAATAATTGGATAGTGGATTTTTGTCTAGGAATAGAG
TTGCTGAAATTAGGAAATTAAAATAACATGTATTTATACTAAAAATTATCCATTACTTATATAAAATTCAGTGTAATTG
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GGTAAGTGTGAAATTGGCAGACAGAGAAGGAAACTGCAAGAAGAGANCATGGATCCTATCAACAGAATCATTCAGCCAT
TCAGACACTTTGTGACAACCTAAAAGAGAGAGGAGGGAGTGAGATTTTTGTATGAAGTAAGAATTCAGTTGTTTGTACAT
GAACAGCATATGCTAGCCTGCTTTGAAGACTGAAGTTCTTGGCTTTCAGTTTATAAACCAGTTCTATCTGGGCAGCTT
GCAGCCAAATTGTGTTGTGGAGGAATGGGACTCAGGAAGCACGGGCACCCTGAAATAGGTGGATGTGGTCTGTGGAATA

Fig. 9.61

GGTGGAAGCACACACTAGGGTTCTACCCTTAAGAAAATGAACCTTTGCTGAGTTATCAAAGTGAGTACTTGCTATTTCT
CAGGATAGCCATGCCACAAACACATATTTACAGTAAAATTCAAGTTGGTAACAATTAATAATGACTGACCTCTAGTTTAT
CTATATATTATTGGAACAAAAGGGCTTACTGAGCAATGCCAAGGAAAGAACTTCCAAGTGGGTCTTCTTACTCTACA
TAAAAGAATAAACTATTTTAAAAGACTTTTGGCAAGTCCACTGTTTATACTACCATAAGTCTTACCTTTCTGTTTTAAA
GCAAGCTTGGCAGGACAGTTACTTGGAATAAGTTGTCAAGTGTGGTGGAGTGGTAGCAGTTGTGTCTGGAATTCCT
CATTCCTTTCTCTCGCTCTCCAATAACCTACCCTGTGAGGCTGTCTCCAACCCAGAGCCCTGACCAAGTGACCATGCTG
CTAAGCTGCTTGATAATAGTTTAATATCAGTAAAGGGACACAAGTTTAAAATTATTTGACATATTAGTGCTATCAGTTA
ATGACTTAAAAATAGCATCTTTGGTTTCTAGGTGTTGACAAAGATTCTCTAACTTACCAAACCTTAGCCATGCTCCTCT
GAACCCCTTCTTGACTAGGCCTTAACCTTCCATATCACAACCTACAGATTCTCAACACCAATGATTTTCATCCACTCGTGCC
CCACATTAAAAGACTTACACAAACACTAGAATAATTTCTAACAGCTCAAGGCCACATCCCTAGGACTACCCTACCCAC
TTAGGATGCCTGCCTGAGGAAGCTCAAGGTTGCCAGGAGAGTTTACTATTTCTTCTAGCCAACACCCTGGAGCTAGGCC
CAACCACCCTTCTTAGAGCATTTACTAAAAAGGGCTTACAATTGTGAATCCTTGCCCTGTAACCTTTGATAAATATAT
GCACTTCTTACTGCTCAAGAGTGTCTTTCTCAAGGACCCAAGAGCCATTCTTCTAAAAAGTAACCATCAGGAGAGAGAG
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TTTGTAATTTTTCACCTCTCTGACTTCACTGAGCCTGCTCTACCCCTTCCCTTTTCTACTCTCTCATTCCTCTTCTGAA
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TGGTTCAAATCTGTCTCTCCCACTTTAACTAGTGTCTGGCTTCCCTTTATCTCTGACAGTGTGAAGAATTTGGAAATTGC
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CTGCTCATACTCCTAGAACAGGCCTTTGGTGTATTTTGCTATATATAGGAGACTGGGACTAGAAGGTCACCCTGTTTCAG
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GCAAAATTACTCTTATGGAAACAGGGGTAGGAAATCTAGAATCGTATTTATTCCTTCTTTTCTCTTTTATACTTATTTGG
GGAGTCTTAGCCTCATTAAGAACAACCTGGGAGGGTAAAACACAAGTTTACAAGAAACTAGTTTCTCCTCTTCAGAGT
TCTTTCATCCTGGCAGCTGAATGGACTCAGCAGCTCTCGGGGGCAGTGGCTTGAGTGTGCTGGGCTTGTGTCTCCTAA
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GCTACTGGGATCACTGCGATGAACCTTTGCCTGAGGTCTGTAATGTGATTTACAATAATTATTCCTTACTCACT
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TTTGCAATATGCCCTTGGATGTGAGGATGCTATACCTGGATTATCCTGGCACAGAGCATGAATTATAATCCTGATCTC
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GACAGAGAGAAGGCTCTTTTTTTTAAAATGACTCATAGCCATATCTATATATAACCACTGAGTGATAAATAACTCATAG
CCATAGCTACATATGGCCACTGAATTGAGTGATTAATAATTCTGTTGTTATAAATTTGAGTGAATATATTTTGGTCAAT
TATGTGGTTGTGGCTATAAAGATAGCTAATAATCTTTATTAGAATTTGGATACTGTTGATTTTTTTTCTTGTAATTTTAT
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CAGCATGGATGCAGCTGGAACCCATTATCATAAATGAATTAATGTGAAAAACAGAAAACCAATNTCCCATCTTTTTCAC
TTATTAGTGGTAGCTGCTTATAAGTAACCTGGACACACACAGACATAAAGATGGAAACAATAGATACTGGGGACTCCGAA
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AAGTATGTTTAAATAGTAGCTGGAAGAGTTATTTAAATCCACACACTAACAGCAACCACAGTGAATACGGTTTAGGTAAT
ATTTTCTTTTATCCAACAGCTAGAGTGAGATGCACAGAACAATGAACAAGCACCTCAAGACCTTGGACTTGTAGAGAT
TCAGAGTGAGTTTCAAAGCAAGAAAAGGCAGGAACATGAGTAGCATTCTATAAAATAGCTACTGAAGCAGAGTAGCTA
AGGAAAGATGATTGTTGTAGAAACACAATCAGTGAAATAGTCTGGTAGAAAGACTATTCCTTAAATTTCTTATACTCCC
ATAGAATTTACTTCTAAACCATACTGCCCTTAGACTTGAGCAANAATATTCCTTGCCAGCCCGCATTTTCAGAGGATC
CTAATTAAACCTCCTCTTGCCTTCCCTTCCCCACAGGGTGAGCATTCGTGAAGTCCAAGTAATCATATCCCTAACC

CTCCCATGTAAGCCATGGTCTGGGTATCTGGACTCTAGAATTTCTGCGCAGGGTAGTCCAAAACCAAGACACAGGTCTA
TCCTTCAGGGGGGAAGTTTGCNTTCCACACAAAGGGAGGCCAAGGGGCAGGGAATGTGAATGAATGAGAAACCATGGCGT
CCATACATAAGTTTGC AAATCTCAGTGAAAGAATAAGTTGGCATGAAGAGAAGGAGGGGTAGGAGAGAGCCTGGGGGCC
AGCTTCTTCTGAATAAGTAAATTCTGGTGCAGGGCTTTTCTTAGGTGCGTTGCATATGTTACCTCATTTAATCCTGCAG
AACACTTCAGGGTAGTGTTCTACTGTCACATTATGGGACAGGGAAATGAGGCTCAGGGGAGTTAAGTAACAGACCCCAAG
GCAACTGACTTGGTAAGAAGTTGAGCTGAGATTAAACCTTAGGTGATTGAGCTCTAAAGTGCATGTATTTTTCTACCTT
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TATTTATTCACCTTTACTATTAAATGGTATATACAATAACAACAGTGGTGTAGGTGAGGAATTTTCATGTCAAATATACTGA
AAGGGACAGTTCACTCCTGATCCATCTGTACTGGATGGATGTGTTGATCTCCCTGTGGCCATGTAGCTTTTGGGCTTAT
CTCTCCATTTGTGCTAGTTTAGTAAATTGCTGCAAATGTTCTGGCTAGCAAGGACTCAAATGTCCCAGAACAACTTTTGC
TCAAAGAAAATATTAATAAACACCATATATGCTTCTGTAGTCCAAAATGTTTTTCTTTTCTTACTTAAAAATGGCACAA
ACTCTGAGTATTAAGGTGATTCAAGAAGTTGTGAGAATGCCTGTTTCTCAGGAGTTGATTGGCAGGGTCCCTCTGTTGG
TGCGGCCCTGCTTGTCTGGGCTGTAGTTCTTCCATGTGTGTGGGCCACCCATGCTGGCTCCTGGANTGGTCAGTGTGCT
GTCCTGGGGCTTGGTCTTTCCCTCCGCTCACCCCCACTTATCATGGATGCCAAGACCATTCTCTGCTGTTGGCAGAAAG
CCATGGAACATACTTCCCTAATTCCTGCTATAGGCTCATTTGGTATTTTCCCATGTGCTCCCATTTGTACACTTAATT
GTTTGGGCTTATTTGTCTGTCTTCTTGTCTAAACTGTAGGCTCATTTGCGCACAGGCTTTGTGTCTCTCTTGTTTACCA
TTATATCATCACATCTAACACAATGCCTGGTACATAGAATACTTAATGAATTTTACAGAATGACAATGGACTGCCATC
ATAAFTGAGTCATTAACTACTTTTTCAA AAAAGTTCTAGCATTAAGGTATCAGATCAAAGTTTCTTTTCACAAAAATCT
TACATTTTCTCTATACTGTAATTGTATCTACCTGCTTGGAGACTCACCAAGAGTGTAGTCACTGTGTACCCTCCCAGA
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TGCTCACTGACACAGGCTTAACTGGTTAAAGAACTCACAGGTTGTTTCTCACCATTAATGGCTATTATGTTATTTTACC
ATCAGTTATACCTCATTTCTGCTTCTATCTTTACAACFGCAGGCTTTGGTCAATTTTTTAAAGTCGTTCAAGTTTAGATTG
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GCTCTCATAGAGTCCCAGGTAGAACTGATGGTGAAATFGACATAAGGTAGATTAATAGGTANATTAGTCCATTCTCATG
CTGCTAATAAAGACATACCTGAGACTGGGTAATGTATAAAGGAAAGAGGTTTAATTGACTCACAGTTCTGAAGGTCTGA
GGAGGACTCAGGAAACCTACAATCATGGTGGAAAGGGGAAGCAAACATGTCTTCTTTACATGGTGGCAGCAAGGAGAAG
TGCTGAGCGAAGCAGGGGAAGTGCCCTTTGTAAAACCATCAGATCTCATGAGAACTCACTCACCATTATAAGAACTGCA
TGGAAGTAACCCCCACCCATGATTCAATGACCTCCTACTGGGTCCCTCTCATGACACATGGGGATTATGGGAACTACA
GTTTAAGATGAGATTTGGGTGGGGACACCCTAAACCACATTAATAGGATAAAAGCATGTAAGTTTACATAACACAGG
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CTGTGAAAAAGTAATAAATTATGTGGGAAGACTAAAGGAAGATCAATAAGAATTATTTTAAACAAGGTCTGTGTGTACA
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GTTCAAACATAATCCTTATACCAAATGGCATATTTCTGGGGTACATATTCTGCCATTCTTCACTACTAAATAACAGAACA
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CCTAAACTGCCATATAGCAAATATATAGTGAAATATTATACAGCAATGAAAATGAAACAGTAGTTTTACTGCTATAGGC
AACAAATATAGATGAATATAATAAACATAATGTTGACTAAATCAATTAAAAAAGAGTACATACTGAATGACTTGATTCCA
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GGTTGTACTTTTTGTGAAAATAGGCTGGTGACAATTATCTGAGGTCAGAGGAAAGATTCTGTCTCATCATCATTTGTCATGAT
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CTTCACACGTAATTTAGTTACTTTTCCACAATTGCCTCTCTTTGTTCAACCAAGTATAACAGCATTTAGGTTTGGCCAC
TTGTTTGGGTCTTCATTTCTTATGAGGGCTCCTGTGTCTATGTAAAAGTTATGTGCTTTTCTCCTATTAATCTATCTTA
TGTCATTTTCATTATCAGGTCTACCTGGGACTCTAAGAGAGCAGAGGTGAAATTTTGCTCTGCTACAGTAGGTTCTAT
TAAACGGTTTTCTATGGCATGGGATATGTAAGCAGGGGATAAAGAAATTGTAATAACTCTCAGACTTCTTAGGGAGTTA
GGAAGCTTCTGAAAAAGACCTATCTAAAAGTAGGAGAGAAAGCCTGGGAACCTTTTACTGTGTCTACCTCAGTGAAAT
ACATAGTTGTTTGTCTCTATAATTTGATTTAATGTCAATGCAATCTAAACTTGAACGACTTTAAAAATGACCGAAGCC

Fig. 9.63

TGCAGTTCTAAAGATAGAAGCAGAAGTAAGGTATAACTGATGGTAAAACAATATAATATCTGTTAATAGTGAGAAACAA
TCTGTGAGTTCTTTAACCAGTTAAACCTGTGTGTCAGTGAGCGAATAGCTCTGTGTGGGCAAAACCATTGCTTAAGAGAA
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TGGAAAAAGCCAATCCCCAAGGGTTGCATACTATATGATTCAAATTATATAACATTGTTGAAATGACAAAATTATGAAA
TGGAAAACAGATTAGTGTAGATTAGTAGTTGCTAGGGATTAAAGAAGGGAAGGAATAGGAGGAACTGGGTGTGGCTATT
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AATTATCTCAAAATAAAAAAATAAAAAATAAAGTACAAGAGAGTCAGGGAAAGTTCTATGTAGTTTCTCTTAGATCAGT
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CCATTAAGGACAATATCAATCATCAAAGACATCTACAAAGTTTCTGAAATTCTTCATGCAGACTTTCAGGGGACCATTA
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GACCAGTCATTAAGCTATTGTTTAGTTCTCATTTGGCTTAGGTTGGTTCAGGGATCAGCTTAGGGAATAATAGAGGCAAT
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TCACTGAAAATAACTGAGTGACAAAGTCATATTTGTTATTTAGACAGCTCTTTTGTAGCTTGGCACATAAGTATTGGTGA
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GT
TTGGTTTTCTGTTCCTACATTTGTTCACTTAGCATAATGGCCTCCAGCTTCAACCATGTTGCTGCAAAGGACATGATCT
CATTTTTTATGGCTGCATAGTATTCCATGGTGTATATATAGCACATTTTCTTTATGCAGTCCACCACTGATGGACATTT
AGGTTGAGTCCATGTCTTTGCTATTGTGAATAGTGCTGTGATGAACATATGCATACATGTGTCTTTATGGTAGAATTAT
TTATATTCCTTTGGGTATAAACCCAATAATGGGGTTGCTGGGTAAATGGTAGTTCTGACTTAATTTCTCGGAGAAATT

Fig. 9.64

ACCAAACCTGCTTTCCACAATGGCTGAACTAAATTACATTCTCACAAGCAGTGTGTAAGCGTTCCTTTTATCTGCAACC
TTGCCAGCAAATTAACAAAAAACAAGTATTTTTTTGACTTTTTTAATCATAGCCATTCTGACTGGTGTGAGATAGTATCT
TATTGTGGTTTTGATTGTCATTGGCCTAATGATTACTGATGTTGAACATTTTTACATATTTGTTGGTTGTGTGTATGTC
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CCTTATAGATGCTGGATATTAGACCTTTGTCAGATGCATAGTTTGCAAATATTTTCTCCCATTTCTGCAGGTTGTCTGTT
TAGTATATTGATAGTTTCTTTTGCTGTGCAAAAGCTCATTAGTTTAATTAGATAACCATTTGTCAATGTTTTATTTTTGT
TGCAATTGCTTTTGGCATCTTTGTTCATGAAATCTTTGCCAAGACCAAAGTCCAGAATGGTATTTTGTGGTTATCTTCC
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GGTCCAGCATCAATCTGCATTTGGCTAGTTAGTTATCCTAGCACCATTTATTGAACAGGGAATCCTTTCCCCATTGCTT
GTCTTTGTGACTTTGTCAAAGATCAGATGATTTTACGTGGGTGGTATTATTTCTGGGCTCTCTAGTCTGTTCCATTGG
TCTATATGTCTGTTTTTGTACCTGTATCATGCTGTTTTTGGTTACTGTTGACTTGTATAGTTTGAAGTCAGATAATACGA
TGCTCTAGCTTTTATTCATTTGCTGAGGATTACCTTAGCTATTCAGGCTCTTTTTTGGTTCCATATGAATTTTAAATG
GGTTTTTCAAATTTTTTGGAAATGTCATTGGTAGTTTGACAGGAATGGCATTGAATCCGTAAATTGCTTTGGGCAATA
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TGCACTTTTCCAACAGTCTTAGCAAATGGCACACCAGGAGATTATATCCCGTGCATGGCTCAGAGGGTCTATGCCCAC
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GCCTGCCTCTGTAGACTACACCTCTGGGGGCAGGGCATAGCCAAACAAAAGGCAGCAGAATCCTCTGCAGACTTAAATG
TCCCTGTCTGACAGCTTTGAAGAGAGTAGTGGTTCTCCAGCACGCAGCTGGAGATCTGAGAACGGACAGACTGCCTCC
GCAACTGGGTCCCTGACCCCCAGTAGCCTAACTGGGAGGTACCCCCAGTAGGGGCAGACTGACACCTCACACGGCTGG
GTACTCCTCTTAGACAAAACCTTCCAGAGGAACGATCAGGCAGCAACATTTGCTGCTCACCAATATCCACTGTTCTGCAG
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CAAAGGTAGATAAAACCACAAAGATAGGGAAAAAACAGAGCAGAAAACTGGAACTAAAAATCAGAGCACCTCTCCTT
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AGCCAAACTAAGCTTCATAAGTGAAGGAGAAATAAAATCCTTTACAGACAAGCAAATGCTGAGAGATTTTGTCAACCACC
AGGGCTGCCCTAAAGAGCTCCTGAAGGAAGCACTAAACATGGAAAGGAAAAACAGTACCAGCCACTGCAAAAACATG
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TAGCAAATGTAAAGAAGCAGAAATTATAACAAACTGTCTTTTACAGACCACAGTGAATCAAATCAGAACTCAGGATTAAG
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AATTGATAGACCACTAGTAAGACTAATAAAGAAGAAAAGAGAGAAGAATCAAATAGATGCAATAAAAAATAATAAAGG
GATATCACCCTGATTCCACAGAAATACAACTACCATTAGAGAATACTATAAACACCTCTATGCAAAATAAACTAGAAA
ATCTAGAAGAAATGGATCAAGTCTGGACAAATACACCTCCCAAGACTAAACCAGGAAGAAGTTGAATCTCTGAATAG

Fig. 9.65

ACCAAAACAGACTCTGAAATTGAGGCAATAATTAATAGCTTAGCAACCAAAAAAGTCCAGGACCAGATGGATTCACA
GCTGAATTCTACCAGAGGTACAAGGAGGAGCTGGTACCATTCCTTCTGAAACTGTTCCAATCAATAGAAAAGAGGGAAT
CCTTCCTAACTCATTTTCTGAGGCCAGCATCATTCTGATACCAAAGCCTGGCAGAGACACAACAAAAAAGAGAATTTT
AGACCAATATCCCTAATGAACATCAATGCAAAAATCCTCAATAAAATATTGGCAAACCGAATCCAGCAGCACATCAAAA
GCTTATCCACCATGATCAAGTCTGCTTCATCCCTGGGATGCAAGGCTGGTTCAACANACGCAAATCAGTAAACATAATC
CAGCATATAAACAGAACCAATGACAAAAACCATATGATTATCTCAATAGATGCAGAAAAGGCCTTTGACAAAATTCAAC
AGCCCTTCATGCTAAAACTCTCAATAAATTAGGTATTGATGGGACGTATCTCAAAATAATAAGAGCTATCTATGACAA
ACCCACAGCCAATATCATACTGAATGGGCAAAAACCTGGAAGCATTCCCTTTGAAAACCTGGCACAAGACAGAGGGATGCC
CTCTCTCACCCTCCTACTCAACATAGTGTAGAGTTCTGGCCAGGACAATCAGGCAGGAGAAAGAAATAAAGGGTAT
TCAATTAGGAAAACAGGAAATCAAATTGTCTTTGTTTCCAGATGACATGATTGTATATCTAGAAAACCCCATCGTCTCA
GCCCCAAATCTCCTTAAGCTGATANGCAACTTCAGCAAAGTCTCAGGATACAAAATCAATGTGCAAAAATCACAAGCAT
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GGGGAAAGGATTCCCTATTTAATAAATGGCACTGGGAAACCTGGCTAGCCATATGTAGAAAGCTGAAACTGGATCCCTT
CCTTGCACCTTATACTAAAATTAATTCAAGGTGGATTAAAGACTTAAATGTTAGACCTAAAACCATAAAAAACCCAGAA
GAAAACCTAGGCAATACCATTTCAGGACATAGGCATGGACAAGGACTTCATGTCTAAAACACCAAAAGCAATGGCAACAA
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GCAACCTACAGAATGGGAGAAAATTTTTGCCATCTACTCATCTGACAAAGGGCTAATATCCAGAATCTACAATGAACTC
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GAGATACCATCTCACACCAGTTAGAATGGTGATCATGAAAAAGTCAGGAAACAACAGGTGCTGGAGAGGATGTGGAGAA
ATAGGACCCTTTTACACTGTTGTTGGGATTGTAAACTGGTTCAACCATTGTGGAAGACAGTGTGGTGATTCTCTCAGGG
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AAAGACACATGCACACGTAFTTTATTGAGGCACTATTCACAATAGCAAAGACTTGGAACCAAGCCAAATGTCCAACAA
TGATAGACTGGATTAAGAAAATGTGGCACATATACCCATGGAATTTCTATGCAGCCATAAGAAATGATGAGTTTATGTC
CTTTGTAGGGACATGGATGAAGCTGGAAACCATCATTCTCAGCAAACCTATCACAAAGACAAAAACCCAAACACCGCATG
TTGTCACTCATAGGTGGGAACCTGAACAATGAGAACACATGGACACAGGAAGGGAAACATCACACACTGGGGCCTGTTGT
GGGGTGGGGGTGTGGGGGAGGGATAGCATTAGGAGATATACCTAATGTTAAATGATGAGTTAATGGATGCAGCATACCA
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CAGTTGGGCATGGCAGCATGTTCCCTGTAATCCCAGCTACTCGGGAGGCTGAGACAGGAGAATTACTTGAACCTCAGGAGA
TGGAGGTTGCAGTGAGCTGAGATCGTGCAACTGCATTCCAGCCTGGGTGACAGAGCAAGACTCCATCTCAAAAAA
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TTTGCTAGTATTTTGTGAGGATTTTTCCACCTATGCTCATCAAGGATATTGGCCTGATGTTTTCTTTCCTTCTTCTGT
TTCTTCCAGGTTTTTGGTATCAGAATGATGCTGGCCTCATAAAATGAGTTAGGGAGGAGTCCCTCCTAGTCAGTTTTTTT
GAATAGTTTCAGATAGAATTGTACCAGTCAGCTCTTTGTATGTCTGGTAGAATATGCCTGTGAATCTTCTAGTCCTGG
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TTGTTTTATTGGATCATCTCTCTTTTTTTTTCTTTGTTTGTCTAGCTAGTGGTGTATCAATTTTATTATTCTTTCAA
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GTTATTTCTCTTCTCTGCTAGCTTTTGGGTTGGTTTGCTCTTTTTCTGTTTCTCTAGGTGTGATGTTATGTTGTAA
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TTATAAGCATCTGGCATTCTCTCTGCTTGCCTCACTCCATCCTGCCGCTTGTGAAGAAGGTGCCTGCCTTTCTTTG
CTTTCTGCCATGATTGTAAGTTTCTGAGGCCTCCTAGGAGTGCAGAACTGTGAGTCAATGAACTTTCTTCTCTTTATA
AATTACCCAGTCTCAGATATTTCTTCATAGCTGTGTGAGAATAAACTAATACTACTGATTTCTATTTTCATTAAGCTGT

Fig. 9.66

GGTCTGAAAGTGTGGTTGGTATGATTTCCGTTTTTTTTGAACTTGCTAAGAATTGTTTTATGACACATTGTGTGGTTGAT
TCTAGAGTATGTACCATGTGCAGATGAGAATAATGTATATTCTGTATTATTTTTGGTTGGAGAGTTCTGTAGATGTCTGTT
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TGTGAATAGGGTGTGAAGTCTCCCACTATTATTATGTGGTTATCTAAATCTCTTTGTAGGTCTCTAAGAACTTGT
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GGCAATTTACAGTTCCATTGGCGTGTCTATGGGCCATTTCACTTTTCAAGATGGTTTGTCTGGCTGAATTTAGAGGCTA

Fig. 9.67

Fig. 9.68

CAAAACCCTAAGCTAGCAGAAGACAAGAATAACCAAGATCAGTGTTGAATTGAAGGGGATAGAGACACAAAAAATCCT
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TTTGCTCAAGTAGAGGGATTATCTGAAAGTACAAAAGAATTTGATAATTGAGAAATACATTATTTTGAATGCACATTT
GAAGTAGCTAAAAATTCTACGCCACAGCTGAATGCACTGGCAGTTTGTTAATCACATCCATAAACATGAGTCTTCTACA

Fig. 9.69

ATTGGAGTGGGCCCCACACTTGGGAACATCTTGAAGATGTTTCTTCTTTGGCTGTGCTAAATTAAGCATGCAAAGAAGTT
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TAAAGTGTATTTTCAGTAATTGGAAGTGTGTTTTGATTATACAATTGGATAGAATGTATAATAAAATGAAATGAAAGAAA
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CATAACATACTTCTCGATATATTGAATTCCATGTAAATGCTTATTTCTTCAAACCTTTAAATTTTCTCAGAGTATTTTCTC
CCTTCTGTTATTAATCAGTTCTATTCATAGTGGATCTTAGAAAATTATCCAGTGATTAATATTTCTCCATATTTGTAGC
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AGTGGCGCCGTCTCTGCTCACTGCAACCTCCGCCTCTCAGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTAGCTG
GGACTATAGGCGTGTGCCACCATGCTAGGCGAATTTTTTTGTATTTTTTAGTAGAGATGGGGTTTTACCGTGTTAGCCAGG
ATGGTCTTGATCTCCTGACCTTGTGATCGCCTCCCAAATGCTGGGATTACAGGCATGAGTGCCCGGCCGCGTTACATC
TTTTTAAGTATTTTAAGTAAATTGGTAGTCAGTGTTTCAAGGATCCTTCTAATGACTCTTCAACAGGGGTGGCTTATGAA
ATAGTTCATCAATTTTTTTTTTCTGATAAGGGGAGGTTGAGGCAGGAGGATTGCTTGAACCTTGGGGGGGCAGAGGTTTCA

Fig. 9.70

GTGAGCCGAGATCATAACACTTCACTCTAGCGTGGGTGAAAGAGTGAACTGTCTCAAAAAAAAAAGGGGGAATTTAC
ACTCTTGAAAATAAATATTTTCATAAGGAACTTTAGAGTTCTCCAATATACATGATCAAACAAGGACCTGTTACCATTT
TGGACCATGTCATAATAGAGAGGAGATAGTTCAAATTAGTCATATGTTCCAACAACATGATTATGGAACTAGTC
TTAGAATTGAGAGTAATTGAAGTTTTTGTGTTTTACTGGTATTAATAAATGGTATTACTCAATTCGTTGACAGTACCAGT
CTCTGCAATATCTTTTGTGTTGGGGAAGGGGGAGAGGACCTGTTTCGTCTAATTAGAAACACATCTACATTTTAAGAATAAA
ATATTTTACATATACTTTTTGTGTTATTAAATCTGCAAACTCTAGAATTGGAAGAAATAGCCTTCATAACTCTTCACTGCA
AAAGTTATGAATGTTTGATAGAATTAATTAAAGCATTTCAGTAGAATTAGACTTGTTTGGGAAGGACTGTAGGATCTTTGG
CAAGAAGTGTGTTTTATATTGTTTCAGATGTATACCATTTTTCTCTTAAGGTTTACAAGTTAATCAATAAAGATTCTTGG
CAGAGCTAAGTACAAAAGAACAATATGTATTTTCGCATACCAAATGGATCTAAGTCTTAAGTGTTATTTGATGTCTCGAA
ATGTTTGGCTTTCAACTTTGTTTAAATGAATAGTGTGTATACAGTGAAGACAGGCTTTACTTAGCCATGCCTAGACCCT
CTGTGGATTCTCTCATAATCCTCAGTTATTGTAAACCATACTTAGTGAGACCAAAGGATATTTTGTGTTTGGCTGCATG
GTATTATTGGAAAACGGTAAAAATTTTATTCTTTAAATAATGTGTTTTCTATTTCAGAAAAATAATGTCTATAGAGATA
GTTATTTCAAATGTAGTTTTCAATATGCACTTGTGGGTATGAACATGAACAAACACATGCAGACACACAGAGTCTGAC
GTTATATCTAGAATGTTGAATCCTACTTTTCTACTTAACAGGAATGTTTCCCTATGTAACGAAAAAGTCTGTATAATGT
GTCTCTATTATGTGTATTATATGAATTATATATCCTAATTTACTTAACCATTTCTTTTATTGTATTGGTCATTTTCCCAT
ATTGCATACAAATTATTAGGGAACATCTGTGTGCAAACATTTTTATTACATTTTGGATTATTTACTTAGAATAGATTC
CCGGAAGTGAACTACCAGGTCAAAGGCTTTTCCCAATTTATTTTTCAAAGAAGAAATCCCACTGTATTCTATACAAGT
ATGCTTGTCTCACTGCCAGTTGCTAGCCTTACATGGTTTTTTTTTTTAAATTTTGTGTTTTTATAAATAGAATCTCATTGT
TTTAGTTTGTATCTTTTGGATTACTAGTGAGTTTGAAATTTAAAAACCTGTCTTTTCAGGCCCTTGTCTTTTAGGGTTT
TTAGGATTTTCTTAACGATTAGCATCATCTCTCTCTAAAAACGTTATATTTGTTTCGTTATATTGTAATAGGTTTTTTTT
TTGCTTTTAGCTATTAAGCTTTTAATTATATACTTGATTTTTTAATTTAGAAAAGTAAAATTTTCTTCAATGGAGTTCTT
TTTTAAATTGAATTCCTTGACATTTTTTATTATTTTGTTCAAAGAACCCTATGGGCAAGTGGAATCTTACACTTTTATTA
CCTGGATAGCGGATAACATTCATAGTTTGATATTTAATCTGTAGCATATAGTTGACTTTTATTTTTTGAATATCCTTGC
TTTAAACTACAACTAAAGGAAGGCAGATGGTTAGCTTGTGTTTCTCATTTCTGAAAATGTCCAAGATTGGAACCAATA
TTATCAGTCTGTAATGGAGGTTGGCAATGTCAAGATGGTTTGTATCCATGTCAAGATGTTGAATCAGTGTCCAGGTCAT
TGCTCATTAAAGATGTTGCTCTTTGGATAAACAGGAAACCCAGAAAAAGATTCTAACAATTTTGTATGCTACCTCACA
TTCCTTCGAGACAGTGCAGAACATTTGCTTCTTTAGAAAATATTCAGGTCTCTTTCTTTGTGTTACAGCACTTACTATT
TGCTCATTTTGTATTATTGGCTTGTGACCCTTTTTTGGGGGGTTACAAATACACTAGGGGAAAGAGTATTTTTTAAATC
ATTATGCCTTTGCTTTATCTATGTCTGTATACTCTCCTATTTGTTATTATTATTTTTTATTTTTTGAGACAGGGTCTCACT
CTGATTGCCTAGGCTGCAGTGCAGTGGTGCAATCACAGCTCACTGCAGCCTCCATTTCCCAAGCTCAGGTGATTCTCC
CACCTCAGCCTCCTGAGTAGCTGGGACCACAGGTGTGCGCCACCATGCACAGCTAACTTTTTGTATTTTTTAGTGGAAT
GGGATTTCACTATGTTGCCCAGGCTGGTCTCAAACCTCCTAAGCTCAAGCAATCTGCCCACCTCGGCCTCCTAAAGTGCT
TGGATTACAGGTGTGAGCCACCCTCCCTGCCTCCTATGATTATTCTATATGTATTTTTTTCATATTAATGTATAAGTT
TTTTCAAGTATGGGTATTCCATTTGAACAAACCATTAAATCTCTGCTCGTCTATAGTCATTTCTTATTAGATGTGTCA
GTTCTTAATTTTATCACACATTAATTTCTTTTTTTGTTCCCTTATACATTTTGTGTTTCCCTGAGATCTTCAGAATGTAAAT
CAATCTCCATTCTGTTATGATTTATAATCATTTACTTCTCAGATGATTCCCTCCTTTAGTGCTTTGTGCCTTCTGTTATA
TTCTTATTACTTTTGATCTTTTATTGTTCTAATTTACCTCTTCTCAACCCATCCCCACACTGGCAACTGTTATATGAA
TGGCTGGTCAGATGGAATCCACATATGTTCCGTATGCATGTATAGATTTTACTCAAGCCACTAGTGGAGACACTATTCC
TACTTACACATATGAGTACCAACCTCTCTGTGATGCAGCAATCATATCTTGACATATCTAGAAAGTTTCATCCTCATCC
ATAAATTACTTCATCTTTCTCCAAATAACTCAGGGAATTTTACATTAACCTACCCAATTTAACAATTTAGGAAAAATCA
ATTCAGTGACATGAGGTAAAAGATGCGCTGTCAAAATTGCAGTGTCTCTAAGGTGATTCTTTTTTGACCAGACTAGAAAC
ACTAATCTGGCACTATTTTAGTTGGGGTTTCTGGTACCTTTCCATGCCATAAAAATCTCTATAAAACAGAACAAATTG
CACATTTAATTATAGTTTCAAAAATTTGCATTCTTGGATGAACTTTTTCTCTTGTGCCACTCTTGCCATAATTGCTAC
ATTGTCTACTGTTAATTTGCATTTTAACTGAGCTTTAAACATTTAACCAGAGGATGGTCCAATCGGAGATTCTATAACC
ATCTCAGTAATTGGTAGGCAAGTTATTTTTTATTGCTATTTTCAGCAGTAGGGAAGTCTCCTCTACAGGGGTGGGGAAAG
AGTTGATAACCTTCAAAACAGACTCTTTACTTTGGGGCTGTGATGAGAGAAACAGAACAGAAGTTAGGAATGGTGAAAT
TAGAAGAGACAGATGGTGAGAGATTATCATGATTTATAGGATCACCCAAATTGTCTAAAATTTACTCCAGAACTTTTTTA
CCAGTTAGGGGCAAAATGTCATGCTTATTTTGAGCTCCTATAGATACATTTAAATCAGACATACTGGACTTTGCATTCTG
TTATTTTTTTTGAGGGGGGTGATATTTAAGGAGTTCTAGAAACAGCCTACATTAGACTTAGTGTTTCCAGCAACTAACAA
ATTTAAATGTTATAAACATTTTTCTAACGCATTTTCTTTTAAACCCTCTGGGGAAAGACATGAATGTAAGATGACTGA
TTCTCAGATTATCTGCTTTTAGAACAACCTTCTTTTAGCTGCAAAAACGTAGTGACGGTGTTCTCAATGTGAGACGGGG
AAGATTTTAGTGGGCACAGCAGCTGTTGACTCAAATATAGTTATACATGTTTCTTCTGTGTTTTCTTGTAGGTGAATCA
GCAGTTTAGTTTATTATGAAAATTAGCAATCTATAAGATGATTCTGAAAAATGAATTTCTCAGAAATTTAAGGTACACAC
ATGGCATGATTTGGTGAATATGAAGTGCTGAAGTGAAATACTCATATCCCATTTGTGATCATTGATGTTCTTGTGCAAT
TATGGTCATCATGTCAAGAACAAAATCAATTTGATGCGATTGAAAATAACAAAAATGCTTTATGTCTTTTAGAGAAAAT
ATAAATGATTCATTGTAAATGAGAAACCTCAAACCTGAAAGCACATAAAAATACCAGGTATGAGTGTGCAATTTTGAAGTC
TCTGACTACATGAGAGTTGTCTGTGAAATAAAAATATCCAGTTACACAAAATATTTCTTTATAGTTAAAAAACTTTTTTTC
ATACATATCACTAGTTAACTATTTTTTAGTCCCATCTTTCTTATGTGCCCAGCAGTGAGCTAAAAGCTGTGCATTTAATG
CATTCCTTCATTGTATTTTTTGTGTTTTGTTTTATTTTGTATTGTTGTAAGATGGGAAAGACCAAATGAAAAGATGTCAAAC
TTTCTGGGGGTGGAGAGGACATTTGCATTTTGAATTTAGAAACCTAAATTAGAGGTTTAGGAAGACTCATAAATATGTA
GCAGAGGTTTGGTGGTCACAGCTAGAAGCTGACCACCACCATGGGGCAAATAATGATTTAACCTCCTCCTGTAAGTATT

Fig. 9.71

TGATGTAAGATCATCGGTATATTTTCATCTGTGACATTATCCAAGAAGAAGATGAATCTGAGGGAAACAATGCTTGCAGATTTGAGGGGGAAAAATGCTTGGAGTAGTTAGTCTCGGGATAGCCCTAGGTGATGATAATCCTTGCACAAACACAGCACA CTGTTAGAACAGACAAAAGTCTGCAGAGGTTAAGAAGACCCCTTGTTACATTGGAATCTTTCAATCACTTCTAGAAATAT GATTTAGTTAAATCAATATTTTATTTAGCTCTAGGTACAATATTGATGCTAAAGAAGTTCTTGTTGAACCAACAAATGG ATGCTCCTTTTAAAATGCTGGAGGCATCATAAATGGCATGTTAGCATCAGGTACCTCCCTCTTGGAGCTTACACATAAAA ACAAATGTACACAAATAGCAATAATCAAATAAAATATTAGGGGATTGTGTCAAATGAGAGCTAGTGACAACTGTTGATT TAAAGATTGAGCCATTTCAAAGTTTTAGAAAGTTTTAGGGATATAAACTAAAGAGTCCAGAGAAAGCAGGGAAAGTATAA ACTATTTCTGCCAAGGATGGGGATTTTGAGGCAAGGTTGGATATTGCTATTAGGAAGAAGTGGCATTGAGCTGGGTAG GATTTTTTACATTGAGAAATGTGAATAGGCACGTGTAGAGAAAAGGAGGCAGAGGATTCTGGATTGAGGGGAATTTATTAA TTCTTCATTCTACATTTGCTGATTACTTCTATATACAAGATACTAACTAAGTGCAAGGCACACAGGGTAAATAAGAA TTTTAGTTCTTGTCTTTGAAGAGCCTTCCCGCAGTAGAGGAAACACGTA CTTACAAGAGGCAAGCAATGGTGCCCTAAG GCCTGAAGGGGGCGCTCTCCTACACAGCAATACGAGGGTGGAGTTTATTTTCATGGCACCTCAACCCATCTGACAGCT AGGGGAGCTGCATTTTCCAACAGAAGGTGCATTTTGTATATTTGTAGGCTGTTTTATACACTGCTTTGTGCTGAAGTTA TTCAGGCATTTCTCTTCCCTCTAGACCTTAAGTTCATGGAGGCCACAAGCTAGGTTTTACCATTGTTGTATCCGTCTGT GCTGTCTAGAGTAGTGCTTGGACACACAGTAAGGATTAAAGACTTTCCGAATATAATTAAATTTTGCTTTTACTGGGA AATAAAATATATCTCATAGAAGGCATGGTTGTCTATTCTTCATCTAGTCTTTCAAAGTCTTTACAGTGTGTTGTGTGG CAGAGGCCAACAGTAGTTAATACTGTGTTGAAGCCATACCTTTCATACCTTGATGTTATTTTTGAACAATGAAAACCTTG CAGGAAGTTCTCATCCCTTTGTGGCTAAGATATTTTCCATTAAGGGGGAAACAAAAGAGAAACATTAGCAAGCTACTC ATTTATTTCTTTTAAAGTTACTTGTAGAAAGTCACCTTTCACCCCAATTTACATTGAATTTAAGTTCATTCTACCT ATCTGACAGTAAGTCAATATTCTGAGATTGTTTCGGAGGCTTCTTATTTTTCTCAGCTGTAACTGAATTACCTTTTA AACGATTTTCTAGCTAACAGTAAATGGTCCATAAAACATATGAATAAAAATGAAGGCAACATTTTCATGCTTAAATAGC TATAATGGTAGTCGGTGTAAATGGGCTGGAGTTTCAAAAATAACCAGTGATGAATCTCTTAGCGGCCACATCATATTATC TGTGGTTTTTGCAGAAGTGTCATTTGCAGCTATTCTGACACACATCTTACTCTAGTTGGCAATTTATGATGTTTTATT C TATGTAAACAAATGTCTGTCTATCTTTAGAGTACAAATATCTACGGAGATCAAATGTAAGAAAAAGTGGCCCTGACTT GGTTCATTGTTGTCAAAAATGGTCTTTAAGAATGTTGTTTTCTTTAGACAATATTACTTAAGTATCTACAATGGCTT GTTGTATTTTCATGGGAATGGTATTACTTTACAGCTTCTGCCAGGGTATGTATCAATCTTCTCTCTACTAAGGAAACAG ACAAAATCTGCTATTTCTTAAACAGTTTTTTGTGGCACTTATAAGAATTATCATGTTTCCCGGGTTGTTGTGACAGACT GAGGATAGCTATGCCTGAATTCATGGTGACGGTAAAGAGAACGTGTAGTGTTAAACGGGCGTCTGTTTGCCTGAAGTT GTCCAATCAAAATGTCTTCATTGATACCAGCTATTTTCTTAAAGCTCTGTATTTGCTCAGAGGCACTGAAATGTTCTT TCCCTTTCTTGTGCACTCTGAAACATTTTAGAATGCTTTTCAAATCTTGAAATCTGGTGATTGCATTTGAAACAGTT TTATAAACATGCAAACCCACTCATGTGATCTGCTGGGTTTTCTGTTGAAACTGCCACTCACATGCCAGGGTTTGTACAAA TAGACCTGAAAGGAATCTCAAGGTCATTTTATTGCAATCCATATTTGATGAATTTGGGAAGTTGTGGGCACCACGTCA CTCTGTGAACTTGTAGCAGGGAGCTGAGGCTGGTAAGGTAGTATCTCTTTATTTTCACTTTAGTAGTGTTATATTCACA CATAGCTTTTCTTCTTAGGGTAGAAGTCTTCTCTGCTAACCTTTGATTTTTTTTGAAATTTCAATTTTTTTAATTCTTCA ACTCCCAAACAGATTCGTTGTTTTTCTTCCCTGTAATTTTTTATTTATATCTGTGTGTTTTGATTAAGGTTACTTTCTA GTTGTAATGGGGAGCATTAGGCAGGTTTTTTTTTTTGTGTTCTTTTCTAAATAATCAGGACTTTACACAAAGTTACAA GCTCAATAAGCAACAGCCGAATCCTTATTCTACATATTTTTTCAGCAGAGTGCCCTTATCAGACACTATGTCCCTCTCTAAA GTCTGCACACCAATGTCTATGACATTCCTAGACAGCTATTCCTTAGTACACCTTTGTACTTCAGGTCCCCTTTGTGGCGG TGCATGTTTGGCAGGAATCCAAAATCTGTTAATGACTGTTGTACTTGCTATTATATTATATCATATAATTATTATGTAG ACTGATGTAACATATAAGTAAAGAGAAGTGTTTATGTGAAAACATAAGATGCTAGTAAATTTGCTAAAAAAATTAAC TGTAGAATTAGATGTAGGTGAGCCAATCATAAAAGATTGGGAGGAAATGTGAATGATTCTTAGATTGCTTCTCAAGTGC CTTAACTTCTCAATTACAAAGAAACACTGTAAGCCATAGATGATGCATAATGGATGTTCTTGATGTAAGACATTGTAGA AATCTAACTAAGAGATTCAAACCTCAAAGCAAAGGCCTTGGCTCTACATCAAAGAGTAGCCAACCTATGTGCATTTAAGT GTTGCCATTTATAAAGAATACTTGAGGTATTATTTCTGAAGATTCTTGACTTTAATATATTCAATTTAACAAACTGGCCA ACTACCTATCCTGAATATGTATATGAGAGGGCTTCTAACATGAGAATAAATCACAAGCCTCTAGCTGTTCTCTATTTTT AAAGTGGGGATGAAAGGTGAACAAAGTGTTACAGATTCTCACTATTTGAGTATCTAATAGTGATGGGGAGGCTGTCCTG TTGCCTTAGTTGTCCTGGAGAAATATCATCGGGGCCTCTTTTCTGTGATGCAGCTCATGGCAGAGTACACCACTGTCAT CCTAAACTTTTAGCTAAAAGCAGATAACACACTTCTTTTTCATATAATGCATTTGTATCTGAATTAGGACTTTAGTGTT ACGGTTAAGACCTACAGGCATTGATTACTTTGGGGTGAAGTCTGGTGACCAAGACAGTGTTCTTAAAGTGCAACTT CCTGGGAGTTTCCACACCTAGCTAGGAGATTGTCTCAGGGACTTTTTACCCAGAAGATACTCTATTATTGGTAGGCTT AATAATAGCAGAAATACAGGCTACCTTATTTTCATGATTATGCATTTTAAACATTAATTTTTAATTTCTTGGAGATCAG GTAATAGAAACATTAATAGCTCTCTATACTACCAGGCATAGTTACCTAAACAAGGTGAGTGCTAAATAGGTGTAAAAA TAATGATCAAGCTCCCAAAGTGTAATTTAGTTATTTTGCATGACAATTTTAACAGAAATTTGTCTCCTATCACAAAT TGCAGTTTTACCATATCAATTAGTTGGATCTTGTCTCATCTTTCCGTTCACTGTGCTACCTAGTGTTGGATGATTCTGCG TATTTTAATGTTGAGATGTGACCACACTGTATTCAATCAGCTGGTTTTCTCAGAACTCGCTGAATATGGGACAGTTTTTT TTTTCCCAGTAGTGAGCTTGACTCTGTGCATAGGAAATACACGCAGTCTCATGTGCCTTTCCCTTTTCTGCAGATGG TAGTTTCATGAGCCTCCCTCAAATATAACAGCAGCTCATAGGTGAATTTATCAAAGAGTATGGCCACCTTGGGAGACCTG GCACATTTTACAGGCCCTGCTGTGATATAGAACAATCAACTCAATTTTTTTATGTGTTCTTTCCATTATTGTAACCTCCC ATGAGGTAAAGTGTTAAGCTACATTAAGCCTCATTAAGAGAAATTTGAGTTCATGATGCATATGTAACCTTCGCAGTAAT TGTAATTAGTACAATCTGCTTCTTGAATTTGCTAACATATTTGGGAAAAATGTTACCATTTTTTTATTTCTTTTAAACA TTTAATGCCAAAACTTTTCAGTAGAGCATTAGTACCTCTCACAAAATTTTATAAGTGCTTTGATTGTGTGCAGAAT

Fig. 9.72

Fig. 9.73

CCTAGCCTAAACAACAGAAAAAATAAATGGGAAAAACAAAACAAAACAAAGAGCCTAATGGGCCCATGTAATT
ATAACAAAAGATCTAGCATTCATGTCATCAGAGTACAGAAAGAGACGAGAAAGAGGGATGGGATGAAAACTACTTGAA
GAAGTAATGGTCCCAAACCTCCCAAATTTGGTAAAACACATAAACCTGAGTGAACCATAAACAGGATAAACCCCAAAGAA
ATTCATACCATTTCATAATTAACTTTCAAAAATGAAAGACACAAAGAAAATCTTGAAAGCAGCCAGAGAAAAATTATT
CCTTACTTATATAAGAAAATCAATGAAATGGCAGTAGATTTCTCATCAGAAACCATGGAGACCAGAAAGAAGTGACACA
CATTTTTTTTTTATGTGCTGAAAGATAAGAATTGTGACCTATACTGAGTGAAAACACCCTCTTTAGGAATGAAGAGGAAA
TCAAGACATACTTAGATAAAAAACAGATTATCACCAGGAGATCTGCTGTAAAAGAATGGCTAAAGGAAGTTAGCTAAGCA
GAAAGGAAGTAACATAAAAAGGAACCTTGGAACATCAGGGAGGACAAAAGAACATGGTAAGCAAAAATATGTATAAATA
CCATAGACTTTCCTTCTCCTGAGTTTTCTAAATTATGTTTGATGGTTGACAAAAATTATAACATTTTCTGGTGTGGTTC
TAAGTGTATGTAGAGAAAATATTTAAGGGAATTATAAGTGTGGAGGGTTAAAGGGACGTTCAAAGAGGTAAGGTTTCTA
TACTTCACTTTAAGTGATAAAATGACAAAACCAATAGACTTTGATACATTAAACAAATATGATGTAATACCTAGAGCAG
CTACTAAAAAAGTTGTACAAAACAAAAGTAAAACAGTAGAGTTAAGCCCTAGCATACCAATTATTGCAGTAGCTGTAAA
TAGTCCAAATGCACCAATTAAAAACAGATTGTTACCTTTTTTTTGTGGTAAGAAGACATAAAATCTATTCTCTTAGCAA
AATTTTCAGTATATGATGTAATATTATGAACTCCAGCCCTCATGCTAGTACATTACATCTCTAGAATTCTTTATCCTATA
TAACTGCATTATGCTATGGTACACAGGATGGTTTCTTGATTTCTTGTGGTATAGATCATTATTGGTGTAAAGAAATG
CAAATGCTGGTATGTTGGTGTGTATTCTGCAGCCTTACTGAATTCATTTATTAATTCTAACAGATTTTTGTGGAATTT
TGTATCATGTAATCTGCAACTAGGAATAATTTTACTTCTTCTTTTTTGATATGGATATATTTTTAAATCTGTTTCCTCT
CTAGATGCTTTTTGCTAGAACTTTTCACTATTATGTTGACTAGAAGTGGCGAGAGTGAGGATCCTTGTCTTGTACTGGCTT
TTAGAGGAAAATCCGCTTTTCCCCATTGGATATGCTAATTTTCTTTTGTGGTACTGTAGTAATCATTTTATTATGTATA
TGTATATAAAATATTGTTTTATATACCTTAAATGTATACAATTAAAAAATAAAAAATAAAGGCACATGATGTAAAGGAAA
AAACAAACAGAAAGAACTTAAAGCAAAACAAAGCAAAAAACAAAGACTGGCATTATGGAATAAAAAATATGACCTAACT
ATATGCTGTCTAGAAGCAACTCACTTCAAATATAATGATATAGGCAAGTTCAAAATTAAGATAAAAAATGTATATCA
TATAAATATTAATCACAGAAAAGCAGAAATAGTTATACTATCTGATAAGGTAGACCTCAGAGCAAAAAAAAATACTAG
TGACAAAGAGGTACATTATAAAATGGTGTAAAGAAAAACTTTAGACAAATTAAATTTAACAAGTTTAATCAAGCTAA
GAATGATTGCAAAATTAAACAACCCCCAGAACCCAGAAATAGATTCAGAGCAACTCTGGCACTGCTGTGTAGTCAGAGAGA
ATTTGTGGAAGAAAAAAGAAAGTGATGTACAGAAAATAGAAATGGGATACTGAAACACCTGGATTGGTTACAGCTGGG
TGTCTTATTTGAACAAGGTTTGAGTAGTTGGCTGTCTGTGAATGCTAAAGTATGGCTGCTGTGATTGGCTGAGACTCTG
CTACTTACAAGAGTAGGTTGCTGTCTATTTACACACCCTGTTAGGTTACAGTTCCTATATACATATAAACCATCAGGC
CTAACTTAAATGTGTAAGGAGTCAGCTTTAGGCAAGTTTAATTAGGCATGGTAAAAGGGTCACTTCACCAAGAAAATT
TAGCAATCTTAAATGCATATGCAGACAACAGAGCTGAAAATATCCAAGCAAAACCTGATGGAAGTAAAGGAGAAATAG
ACAAATCTTCAATTACAGTTAAGACTCCAGCCCTTCTTTCTCAACAATTGATAGAACAACTAGGCTATAAGTCATCCAG
GATATAGAAAACTCAGCACCATCAACCAACTGGATCTAATCAACACTTAGAGAACATTCTTTCAATAACAGCAAAAT
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AACAAATTAAATCATAACAGAGTGTGTTCTTTGACCACAGTGGAAATCAAACCAGAAATCAATAAGAGAAAGATAATAGA
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TAAAAAAAAGTTAAGCTTAACAAAAGTACAAATGCACTATAAAATTTGTGGGACACAACCTAAAGCCATGCCAAGAAAA
AATGTATAGTACCAAATACATACATTAAAAAAGAGAAAAAGACTCCATCAATAACCTAAACTTTTACCTCAAGAACCTA
GAAAAAAGAGGTAAATTAACCCCAAAGCACACAGAAAGAAAGAATTGGTTTTGTAGGCTGACTGGGGAAAAATAATT
TATACAACCTTGCTTCTGTGTAATAAAAAAACAATGAGTTCAAAAAACACATTGTGGCCTTCAAATGCCAATACCCTAT
TAAATTACCAGAAATTAATAATTAATAATTGATTGATGGTTACTGTAATGATTAAGATAATGGTTACTTGTGGTGACCT
TTTTTCCAATCATAACCAAACATATCTCTGGAAATGAGTCACTAAGGATGATGCAGCATCTTTCTTTCTCTTTTGATTA
ATCAAAATTATATTAAATATATTTTATCTTATATTAAAGAACAGTTTAGTCATGAATTTAAGCTGGTGTAGAGGATAG
CTAAGTTGCTTCTCATAGTCATTTCCCTTACTTATGTAATGAAGAGCAGAAAAATATTTTTACCAGTGGTTCTCAACCTT
GGCTGGAAATCAGAATCACTTAGGGAGCTTAAAAATATACTGATGCCTGGGTCCCATCCAGCGAGATTCTGAATTGTTT
TATGGCAGAAATTCTCAAAATACAGTATTAGCCTCTCAGCAGCAGCAGCAGCACTTGGAACCTTGTTAGAAATGCAAAAT
TCACAAGCCCCATCCCTGATCTAGCAATCTCTGTTTTAACAACCTCCTTCAAGTGATTCTGAGGCAGCAGGTCTCAAGCT
TTAATGTGCATGCACATCTCCAGGGAATCTTAGGAAAATACAGATTTTAATTTGTGGTCTGGATATTACACACTGTCA
CAGATGCTAATACTGCTGGTAGCAAAAATATAATTTGAATTACAAGGGTCCATAGGACATCTGGATATTTGCATTTTAA
AAAATTTTCCAAGTGGGAATGCAGCCAAGGTGAAAACAACCTGGTCTAGATAGCTTTATGGTACACTGCCAATAGCCCAA
GCAATCTGAATGATCTCTGCTTGGTTTTCTGTACCTGAGGTTGTAGAGTCACTGAAGAGCACATACTTCTTGTCTCTT
TAAAGTGATAATCCGGCTGGACATGGTGGCTCATGCCTGTGATCCCAGCACTTTGGGAGGCTGAGGCGGGTGGATCACT
TGAGGTGAGGAGTTTATGACCAGCCTGGCCAACATGGTGAAAACCTGTCTCTACTGAAAATACAAAAATTAGCTGGGCG
TGGTGGCACATGCCGGTAATCTCAGCTACTTGGGAGGCTGAGGCAGGAGAATCGCTTGAACCCGGGAGGTGGAATTGC
AGTGAGTCGAGATTGCACCACTGCACTCTGCACTCCAGCCTGGGTAGCAGAGCAAGACTCCGTCTCAAAACAACAACA
CAACAACAACAACAACAACGACAAGAACCACAAATAACAACAATAAAACCTAAAGTGATGATCACTGATTTTAAG
TGGCTCCTTAGTTGCCTAAGAATCCCAGTTGTGATGGTTTTATCCCTTATTGTCTAGAACTAATGTTGAACACCCTGCT
TTTTAACTTCATCTTGTTTTTCTCTACCCCCATCATCAATATTTGCCTGACTCACCATCTTTCAAGGTTTACCTCTTCA
TACTCAGCTAAAAATTAGCTGATGAGATGCACAAATAATTCCAGTGTATCAGGAGAGGATCAGCTTCTCCTTTTAAAG
GTAACATGGATCCTTTCACCATCTCCTGATGCCCTGAGACCAATGTCTTGAGAGCATCACATCTTGTTCATGCATTCT
GACTGGGCATGCTCATCTATTCCAGTGTTCCTCAAGAAACACTCATGTATTGATCACTCCGTGGCCAGGCCAGATCTCTC
TCTGGCTTGTCACTGAGTGGACATTTTTTTTTTTTGTATTCTCAAAGTCACCTCAAATTCAGTGTGCCAGACTAGAATC

Fig. 9.74

ATCTTCCCTACCATGTCTGCGTCTCCTCCTGCATTCTCTGTTTCAGTGATAAGTAACTTGGGGGGGCATTTTTGATTCTC
CTTCCCTCTTTTCCACCCCAAATGGAACCTTGTTGTTAGGATCTGTGGATTCTATCACTCTTGATTCCACCTACTTTTCA
CCATTTTGCCTCTACTTCTCTGGTGCAGGTCTTCACCATCTCTTGCTTAATTGCTGCTACAGTGTCTTAACCTAATTTCC
CTGCCTTTAGTCTGTCTGTCCATCTCTAATCCACTTTCCATATGGCAGTGAGAATGTTTATTCTTGCCACTTCTTAGAT
CTAAACTCTTTAGTAGCTCTCCATTGCTCTCAGAATTGAGAGTCCCTTATTATGACTTAAAAATTATTAATTATCTGGT
GCTTGCTCACCTTCCAAGGATCTTCTTTTCCACTTGAACCTGGATGCATTTTCCGTATGTACCATTTTCTCTGTAGGTTC
TAAGCCTTTGAAGAGGTGATGCCTCTGCTTGGAGTGTCTTCCATTCTCCCATTTGCCTAGTTCTTATTCTTTTTTTAT
TTGCTAAGACACTACCTCTGAATAGATGACTTCCCTTGACCCTGCCTCTCTCAAGGGTGAGTTACATGCCCTCTTCTTAT
CCTTAGTGTTAAAATTACCACACTGACCTTCAGTTCCTCTTTACTTGTTTATCTTCCCTCAGTAGCTCCTTAAGAAGGG
ATTATCTTCTTCACTCTGCCCTTAGCATCAAGCCTGATGTCTCTATTATAGTCTGTGCTAAATAATTATTAATAACAA
AATACCCAGTTAGTGTAAGGAAAGGAACCTGCCATGGTAAATAAGTGTGAGAGAGGTTTTGAGGTAGTCGTATAACTGT
TAGAAATACACAAATCACATTATAGATAGTGGCTTCTCTGTTTATCACAGTGCTCAGTTTCTTGTTCCAGGATATTTAG
AAAAAGTATACCTAAAATAAGCGGAGAAGTCATTAGGTGTAATGACATATTCCTTCTGCCGAGGTATTATAATATAAT
TTTTAAAGGGCCTCAGTTAACAATTAAATTCTTTATATCCTTGACAGCAGTTCTTTTTTTCTAGGTTTCATTTCTTCCA
GAACTAGATGAGGATTTTTGTCTAACTGATACAATTCCACGTTCTGAGTCCCTTGCTTCTCTCTCTTTCAGTTTCTGGG
TTGTATGTGGTAGATTGCCCAGTCTATACCTAATCAGCACTAGGTTGGAGAAGAATAAACTGCCTTCATGTTGTGGAGC
TGGCAGTCTTCTCTGTTGAGTGATAAACCAATGCCATTCTAAAAGATCTGCAGACCTGTGFGATCTCTTCTTCGAGGGA
GCTGTAAGAACTAACTTTGATGGAAACACCAGTTTTTAAAAGATTGCAGTTCAATTTCCCTACATGCATATTTGCCTAGGGA
AATTTTGTAACTCACTGAAGTTTACTAAATGAACAGCAAATGAATAAGTATACGGGTGATGACAAACACTCCAGCC
CAACATCTCTAGGGCCTTTGCCAATGTTTCACTAGAATTTTGGACCCTCAAGGTCATCTTAGTTTTACCATGTTCTCCT
GATTTGTGTTGTATGGGGTCATCCTCTTTAGCAATCTGCAGAGCATAATTTTAAATGCTGTCTCCAGGTGGAGGGTTGA
TGATCCCTTTATAGACCACATTTCTAGCACAAATTAATGTCACTGTTTCTTCTGTAGGAGGAGCCTGGCACACTGAATAC
CAGTGGAGTCTTGCTCTCTAGGCTTATCTTGGGAGCTAGTATGCTTATTCTTGTTTCTGTCCCTTTCTTATGCACTGC
CACAAAGGAATTGCTCTAGAATCCATTTCTTCTTCTCATCTCCTCCGTTTCTTTGAATCTTATTTTGTAAAGTTCAAG
CTTCATTTCTTGCAATTAGGGAATAGATCATGGTCTCTAGGTGCTTTGTCAATTCTCAATACTGTGTGCCATTTCTTTGA
ATTACTTCATGTTCTTTGGAGCAGAGGTTGTATCTTTTACTTTATATGGAAAAGTATGGCTTAGTGACGCGTGTAACCT
CAGACTAAGGCAATTTAAATGCTGTATTTCAATTTGAAATGACACATGGAGCTGTTTTTACCACATTTCTCACTGTCTTA
TCTGAAAACAGATTAGCGCAGGGAAGTCTTGGAGCTATTAAAACCTTCTCCTCATATTTTCTCTGGCTTTCAGAATGA
CTATATAGGGCCTTTTCTTTGGTGGGGATTACACCTCACTGGCTCTCATGTAACCAGTCAATTTTCTCTCACTTATTA
TGGGACTGTGAAAGAAAGAAATTAGAAGGTGAAATTATAGGATTAATGTTAAAAGGAATTAGTTAGCTTCTTTGTTCTT
TAATTTCCAGTTACAAGAAAAGAAATTAGATCAATTACATTTCTATGGGCTTGATCTCATGTAATAGTAGCTATTACCAA
CAAAAATATAACCAATTTGGACTTTTAAAGAAATGTTTCAACAGCAGCATTGTTTATAATGGCAAAACATTAGTATCAACACA
GGTGTCCAACCACAGGAAAACAGTTAGATAAAATGTAATTAAATCCTTATGATAAAATGTGACCATTAAAGTTGCTGTT
TTTGAAGAATTTTAAATGTGTGAAAATGTGAAAATGTGCTGATGGTATCATCAGCACCTAGTACCAGTGCTTGGCATGT
GAAAACAGCTTCCAATAAATACATATGTAATAGCTTTATCATAACCAGGAAAGGTGGCTATAGAAAATAATAGTAACCTC
GTACTAAAATGAAGAGAACATCAACTTCTCTCTGCTAGATGGCCACTATCTGCATCTTTTATTCTCTTGAGTATTCTG
GAATCAGTAAAAATGGTGTTTTCAAAGAATCTTATCAGAAATTCAGCTACATCTTTGCTTGGCAATCTGAAGTTTTTGG
AAAGAACGTATTTCAATAATAAAGAAATGACATAGAAATTTTACATGGTAAATATTTAAATAAAATATGACATTTTAAA
TTTAAGTCCAGGGTTACTGTTTCAAGAGCTGAATTCAGACATCTGAATGATAAGCGGAGATTTCAATTGAGAAAATCTCTA
CTTTATCATTTCTTACTGGCTTTACTATGTCTATGGAGAGCTCAAAATGTCATCACTGAATTAGTTTTCTAGCTGACTTCC
TCCTGGTGGTGTTTTATATGTTAGTGACATTTGATTTCTGGACTTTTCTTTTTTCCCATAGGTCTTACCTCATTTATTGA
AATAGAATTGATTAAATAACCTATAGAAATAAAAAGCAAAATAAAACAATAGAGATAAAGATGAAAAATGTGGTATAA
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TTTTATGCTTTTAGAAGATATCATTATTAATAACTAATTTATTAGATTTGAACATTCAGAGCAAAAGAGATGTTACACCA
ACGGGTTCCTTGTCTCCTACTTTCTAGTTTATGTAATGTAATGAGTACCCTAGATTATCAGTTGCTTTTTTGGGGAT
ATGTTTTGTATGATAAAAAGAACAGTTCGATCATGACATTTGCAGTTATCTTTCTTGACGTGACCTTCTAGAAATGA
CTCTTCACCTTCTATCTCTTTCCCTGTCTTGGTTCATGGTATACTCATTCTTTCAATTGTGAAAGTAGAAAGTCAGCT
GACTCTTTGATTCTTCTGTCTCCCTCTGAACCACTGACTTATGGCAATTTCCCTTTAGCATGTGTTCCGTATCTGGACC
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CTTTATTAAGATGTAATGACAAATAAATATTGTGTATACTGATGGTATAAACGTGATGTTGTAATACATGTATACATT
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ATCTACTCTTTTAGCAGTTTTTAAAGTGTCCAATACATTTTATTAATTGCAGTCACCATGGTGTACAATAGATCCCTAGT
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GTGGAATGTAGCAGAGCCTAATACCATATAAATACAAAATACAAGAAACATTTATGTTGTTTTTTGTATTTTTTTGAGA
CAGAGTCTTGCTCTGTTGCCCAGGCTGGAGTGCAGTGGCACGATTTCCGGCTCACTGCAACCTCTGCCTCCCGGGTTCAA
GCGATTTCTCTACCTCGGCCTCCCGAGTAGCTGGGACTACAGGCATGTGCCACCACGCCCGAGCTAATTTTTTAAATAT
TTTTAGTAGAGATGGGTTTCAACATGTTGGCCAGGCTGGTCTTGAATCCTGACTTCAGGTGATCCGCCCGCCTCGGCC
TCCCAAAGTGCTGGAATTACAGGTGTGAGCCACCGCGCCCGGCCAAGAAATTTTATGTATATATGAATGTTAGCAGTG

Fig. 9.75

GTTATTTCTGAGTGGCTCAATAACAGCAGATTTTTCTTTTTGCCTACACTTTTCAATTTTTCTACAATAAATTGCATTG
TTTTTATAATCAGAAAACTTCCTCTAATGATTGGTATCTGTAAGAAAGCTTGTAAGTTATTTAGTTTAGTTGGTCTTA
ATATATTTTTGGATCATGGGCCTCTTTGAGGATCTGACAAATGACATGGAACCTTCTCTCAGAAAAATGAACATCTGCCC
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CTCTCATTTAAGAGATGATGGTGGACCCAGAGGATCTGCCTGCTCTCCACAGGAGGCTGGGAGCAGAGGCAAACTAA
TTCTTAGTACTTTAAGCTTTTGCCTTGAACTAGGCTGTCCAGGAGACTTGAGATCACTTTGCAATGCCCTTCATTATG
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TACTTTCTTAAACATTTCTCTCAGATTTGATGATTTTGTGTAAACAGGCATGATGAAGGCTTTTTTTCTTGAGAATG
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TCTAGAATTTTTCATTTGTGCTTGGTGTGTAGCGGTTCTCTCTACCTGGTGCCTGCACAGCCTGCATGATTAGGGCTTG
GAAAGGTCTCATTTCTCTGGTTTTAAGTGATTTTGTCTTTTCATAGCCTTAAGTTAGCTAGTAAGTGAGGGAAAAGTA
CCACTACCAGCACCCAGAGCCAGCCTTTGACTGGGAATGAACGTGGAAGAGTGCCCTTTCTTTGTGCAAACTCCTAC
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CAAACACAGATCTGGAAACAGCTGTCTGATACTTTTTTTTGCAAGTTGTAAAGCCCTCTACAGTCTAATCTCTGTGGAGA
GCCCTGAATCAAAACAGAGGATTGAATCCCTGAATTGAGCAGAGGAATTGTGGAAAAGGTAAGGAAATGAACTTTTTTG
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GAAAAGCAGAAAGAGCACCATAGTCCCTGAACAATCTGGGCAGAAACACATGGGTTTGGGATAATACACGTGCACTTTT
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GACTCGGGGCTCAGCTTGGCAGGAGGTGTGATGTCTCAGAGCAAGTACAGCATTTTTTTGAAGGAGCAAGGTGTTAATGG
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CCAGTACATGTGAAAGGACTTTAGTGCTCCCTCTTTTGTCTCTGGGCTATCTTCTCCAATTCTGATTTGTGATGTA
GATGGCTGTTACCAGCAACAATGAGAGCGTGGGGCCTGGGGAGGCAGAGAGCTTTCATTCTAGTGTTTTTGTGATTGTT
TTGGCTATAGCTAGGCCAAGGTACTGTCTTATTCTGTTTGTATTGCAGTCAAATTAATCAACAGGCATTTTTTCTCAG
CCATCTTCAAGTCGTGTCAGGAAAAATTTGGCCTACATGAGGTTTAGGAACTATCTTTTATTTCCCTTTATTTTCATG
AACATTTGAGCTTGAGGAGAATGTGAGCATTTTTTCAGATCATTTGGGTCTTATAGTTTTCAGGCTTCATATTTTGTGCATT

Fig. 9.76

CCCACTTCTAAAGCTTTGTTTTTCATATATTTGCCCATTCCTCCAATCTAAGTCTCACCATTAAAGTTAAGTTCACA
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TCTGTGTATTGAAGCAACAGAAACCATGAAGCATTTTATTGCAAGAGCAATCAGGCAACTTCCCAGATGTTGCTAAACT
CTGCTTTAACTGTTGTGTAGAGCTAGTTTTAACTGCATCTCCCACTGCTACTGGAGACAATGATTCTAGAGATGTTACT
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GTAATTTTTACCTTCTCCTATTGTATCCTATGTTCAATAAAATAAATATCTCTAGCAAAGAATTCAATATAATAATGTT
TCCAGTCTTCTTCTTATTTTATGTCTTCCCATAGGGTAGACAAAATTCAGTAACAATAAATGATCAAAATATGAGA
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AGGTTGGCAATGACTTTGAAATTTTATCAAGTCTATCTCTTTTCTTTTGGCAAGACTATTCTTATACCTTTCCAGTA
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GAGGGTTTTTCAGAGAAGGACTCTGTTGTTTATTTTGAATTGTTTCTCTCCATAGGGATGATCAACTTTATATAACA
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AATTCATGTTGTGTTGTTGACTTATATTCATTTTAGACCTTCATATTTTTTGCCTACTCTTTGCAATTGCTAGTATTCCC
CTGCTCTGTGTTAAATAATGCTGGTTCTGTCAAGTAAATATCCAAATATGTGATTTTTTTTAAAAATAAAGGTGTGAC
AGGGTCTATGATGTATAGGACATAGGATTAGCATGGATATGACTCAGTATCCTGGGCTCCAACCATGATTCTGTACAG
CTACTGGTATGACTTTGGGCGGGTAATTTACTGAGCTCAGAGGCCTCTGAGCCTCAGTTTATCACTATCAGATGTGTGG
CCAGATTCAGAGCAGCTTTTTCCAATGGGTTTTCCATGGGATGTTAACAGATATAACATGAAAAATGGTCCCATTTAA
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TATCTATGATCAGGAATCTGTGAGTTTCTAAGGTTATAAAGATAACATAAGAATAAATACTCATAGATAACCACAGG
TAGGTTAAAAAGGGCAGAGTACAGCTTTGCAGGTGGTTCTTACTGGCTTCTTTCTTTGGTCTCGATAAATTTGCCTAT
TACCTTGATTCTGTCTGATAATGTGAAATTATTAATTGGAGAAGTTTGTCCAGATTACCAATATATGAGGAATGGACAA

Fig. 9.77

Fig. 9.78

GCTTTGCTTTAGCAGAAGACACATTGGATTGGGCATCAGGGCAAATGGATTCTGGCAAAAGATATGCTCTTGGAACCAC
TTCTTTTCAGCTATTAAGTGGAATGAATAATGCTTCACCTTGCTTAAAGGCAGGGGCTGGACAAGATAGCATCTTTGACTG
AATATGAACAGAACAACGTATTTCTAATTGTGTTCTTTAGGGGCCAGGGTTGCTGGCATAAGCATGAATTGAGCATCTT
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AGAAACATCAGTGTCTTCACTTTGACAGTGCTACATTATCTTTTGGTTATCATCCCTAGCTAAAAATAATTTTAGAAGG
GCATAGAAAGAAGAGGTATATCTAAGATACTCTCTTCTAAGTACCAAGGTTGATTGATGATTCTTTTCAGAAACAAGAA
ACTTTCTTAAAGTATAATATGTTGATGGCTTTGTTGAGTTAGCAATTTATCACACCACATAGTTCTGTGGGTAATTTAT
CATTGAATCAGGTTCCCTTTACTCTGTCCCAATATTATAGGTTAATCTGGGACATGTAGTTTCTGAAATACCGACAGTC
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GTAACCATTATTTGTTATTGTTATCATGTATTATGTAGTACACATAATTGTGTGTGCTCTAATTTATATGAATGACAGC
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CTAAATGTCATTATATGGCACATGATTATACTTGAAAATAATCAAAGTTATCAGATAGAGAAGGCTTAGGGATGCAGAA
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TATAAGTGAAAAAATACCTATGCTGCAAAAAACTCTTTTAAATGTTGACTCCAGTTTCAGCAGAATAAATTTGTCTGATG
ATAGCCCCAGTGTCTACAGAAAATATACATTTTCTATTACTGGGGTTACAGTGTATGTAGAAGCACTGTGAGAAACCTT
GACATTTTGGCAGAAAAGGATTTTTAACAATAAAATTCAGAAAAGGTCATATTAGTAGAAATTTTGGTAGTCATTTTA
AATGTCTCCACATTATATGGCATCATAAATGCATTTAATATCAGATTGAATTATGGCTGTAATCTTTAGCCTTAGTAG
TCTGAACATCATTTTCATACCTAGCTTAATATCCTGGAATGTTATGTGTGGTTTTCTGCTTTATTTAAATAGAGGAGTGA
CAAGGGTGATGAATACATTTTATTTCTGTTTTCTCATCCCATTTTCTATTGCAAATGAATTTTTTTTCAGAAGTAGGTA

Fig. 9.79

TGCCTACCTGTATCTTTGCAAAATGCATTTTCAGAAAATAGGTTTGGAAATATTCATTTAAATCCTATTTCTTGTGACCC
AATGTAAACTCACTGCTGTTAAATCTAAAGTGTGTGAGTTCTTACAGCTGAGCAGTAATTCTACTGACATTTTAATGTC
ATGTCCTCTCCTTAGCAGCCTGTTTGCATATTGCATGCAGGCTACATGTTAGGATTTTTTAAACATGAGGTGTCTTGG
AAAATGATTTTGACACAACCTTGTCTCTTGGACCAACTGTATTTTATAAACATTTTAATGCTTTACTCTTGAATGATCC
ATTTCACTAAGAGTCAGAAAACCTGAGTTTCTGTTCATGCCATGGTGACATGATGCTGACATTGCTGAGCTAGTTACTTAT
TCCTTTTACCTAAATGTTTAATAGAACACACCCAGCCCTGTGTGGGCATGAGAGGGAAATGGGCCCAGAAGTTGCAGTG
GTTGCATTCACAAGCCATTAAGCTGTGGTCAGTTTCTCCCTAAAGCATTTGTACCTCCCTTAGACAACTTCCTCTTTGGG
GAGTTGCAATATCTAGAACTTCAGAGGAAGGGGTCCAGCTGCAGGGGATATGGTTAAAGTTTTTTTCAGAGGAGCCACT
TTGCATGGCAGTGGGCCATTTCTCTGCTATCACTCAGGCCCAGCCTACCCACAAGTTACAGTCTTTGGCAAGAAACAGAA
GGAAACAGTGGTGGCTCTTGGAGCCACATGCTCAATTTTTTGATGGTGCTCTTTGCGCAGTGAATGCCAGCTGCTCCTCA
ATCAGGTATGCTGCTGTTGTTCTGAGAGTGATTAGGAACAAAAGGGATGATGCAGATGTGTGTCAGGCTATGGGAAGTGCA
AATGTATATTTTTGGAGCACTGAATGTAAAGAGATTTAGAAGTTTCTGTTGGAAAGCAGGCTATCGAAGCAGGCAGGC
AAAATGATTTTGACACCAATACCAACCAAGGGCAAGAAGTGGAAGCTTGGAAATATACTTGTTTGGGAATAAGAAATTG
ATAAGACAAATATACTTTTTTAAAAAGAAAGATAAGAGAGGTCATGAAAGAATTGTGAAGGTCCAGTCTGTCTTATTTCC
TTTGCTACCACCTGAGTCTGTTACCATGGTCTCACTCTTGCATGAAAGCAGTCATCTCCTACCTAAGCCCCCTTCTGTTC
ATCTTGCTCCTTGGAGGCTGGCCTCCACATCCTTGAAAATACAAGTCTGATTTTGCCTCTCCTCTGCCCAAACCTGCCA
ACAGCTTCCCAACTCATGAGGAGTGAGATCAATGTCTTGCCATGGCTGACATGGCATTTCTTGGGCCTGGCCCTGACCT
TCTCTGTCTTATCTTGCACCACCTCTTACTGTCACCTCGCTCCCTTCAGGTCCCTCTGTTTCCTTGCTGCTCTTTTATAT
GCCATCCCTCCTCAGGGCCTTTCCACCTTCATTTCTGATTAGAATATTCTTCCCCCAGATATCCGTATTFACACACTTTC
TCACTCCATTATCATCTGTTCAAATAGCACTTAAAAAGAGAGGTTCTGTGGCCATGCTATCTAAAAATAGCACACTCACA
TGCACCCATGACTTTCTCTCCACTTACCCTATCTTCTTCACAGCACGTAGCACCCACCTGTCATATTCTATCAAGTTGAA
CCATATGAAATTGCTTTTGGGTGGATCAAGAACAGCTGAATATTGGCCATTTCAATTTGGTTCAAACATAAATATATTTA
TTAATTTGTTTATTATTGTTATTTCTTCTATACTATGATTGAAACCCTGTGGGTCAAAGGTTTTGTTTTGTTTTGGCA
ACATCTCCAATGTCTAGAGCACAGCAGGAACCTTAGACAAGTTGTATTGAATTAATGGAAAGGAAGAAGAAAAGCCATAT
AACTTGTTGACAGTAAGCAGTGAACACTTATATGAAAAATGATCTCTCTCCCTTTCTCTGTCTCTGTCTTGTGTGTGTG
TGTGAGAGAGAGGGAGAGAGAGAGAGAGGAAGCACTAGAGGACTCTAGAAAAAGGTGATGGGATGGGCTTCAGAGGGAAAT
CCAGAAAATTAAAGGATTTATTTGGATCCATGTAAAGTTTCAAGTGAATGTTTCTAGTTGTAGCTAGGGGCTATGTATGC
ACCTATAATTATATATATTTTTTTCCATGTGGTTTTCTGAAACTGGGACCAAGTGGGATCAAACAAGCTTTGGATAAT
ATAATTGTGAACTGAAAAAGTTTTATAATTGGTAAATTACAAGGATTCAGCTTGCCAATTGTGCTGTAAGTTGATTTTTT
ACTAGAGAAAATAAGTTACTTTTTTACTAACAGTTCTAGAAACATGGCTAATAATCTTACTATCATCCCTGCTACGGATG
CTTAGAAATACTGCATGTCTAATCTTGAAAGGAATAAAATCATTACTATGGGTCAGAAGTAAAGGTTGAATTGAAAAAC
TCCAGGGATAAACACTTGAATTGATACTGTGGCTGCCTTTGGGACAGTATTATTAGTCTCACTAACAAAGGGCTTGAGA
TATGAAATGGAATTTGAGGTTCTGTGGGTGGAACAGAGATCTTCAGTACAAAGTTGTGACCTTGGAATGGAAAGGAC
TGAAAAAAATCATCAGAACAGAGAGACAAGGAAGCTGTGTCTTAGAAAAAGAATGTCTCCTTCAAAAACACATCTCT
ATGGATTTGTACCTCATTTAGGTTTGGGAATTGAATTTGTACTACTTGTGTAGTATGGAAAAATCACAACCTTGAAAAAT
TAACTACAGAGAGCCCAGATTACTGATACTTCTAAAGCACCTGGCAGAATCTGTTACAAACTTCTTAGGAGAAAGATAC
CATTCAACCAAGTTTCACAAGATTCCTAAACATAAATATCTTCTAAAGATTAACCTCACAACCTCAAACCTATAAAACATA
AGAACAAAATCCCCCACAAGTGAGAGTTAGCAAACACAAAGAGCAGGATTAGAATAAGACCTAAAAGTGATGGAACCTAT
CACATAGACTATGAAAAATGAATATTGAAAATGATAGAGACATAAATGAATTATAAATATAAGAAAAAATAAGGTTAAA
AAAAGAACACACTTTATAAAGGTGAAAACATAAAGTAATTGAAATTAATAATTTATCATAGGTTAAACAAATTAGATATA
GCTGGAGAGAAAATTAGTAGACTGTAATATAATTTGAAGAAATCAACCAGAAGGCACTGAAAAGAAAATGATAGAAGAT
ATGAAAAAGAAGTTAAGAGACATGTTGGAGAGCATAAGATTCAATATATGTTTCATAGGCCTTCTAGAAGGAGAGCACA
GGAAGATTTGGGAAGAGGCAGTATCTGACTATAATGCAATTAAATTAATAATTTAAATTAAGACAAACAAAAGAAATC
CCACATGGATTTGAAATTAATCAACACACGTCTAAACAACCTCATGAGTCAAGCAAAAAAATAATGAAAATTAGAAAAT
ATTTAAACTGAATGTTAATATTAATTAATTAATTCATCCATTAAATATTGTATGAGAAGCAATGAAAGTTATAGTTAGAAA
GTTATAGCTTTAAATGCCTATATAAGAAAATGAAAAAAGGTACAAATTAAAAAAATAAATTTTAAATTTTTTTGGCTA
CATAGTAGATGTATATATTTATGGGGTACATGAGATATTTTGTACAGGCATGGGGTACAGGCATGCAATGTATCATAAT
CACATGGGGTGTAAATGGGGTGTCTGTACCTCAAGCATTTATTTTTTGTGTACAAACAATCCAATTATACTCTTTTA
GTTATTTAAACTGTACAATTGAATTATTTTTGACTATAGTCACCCTGTTGGGCTAGCAAACCTTAGGTCTTATTCATT
CTTTCTATTTTTTTTTGTACTCATTAAACCATCCCCACTTCCCTCCCTCCTCCACTACCCTTACAAGCCTATGGTAACCAT
CCCTTTACCATCTATCTCCATGAGTTCAATTGTTTTAATTTTTTAGCTCCCAATAAGTGAAAACATGCAAAGTTTGT
CTTTCTGTGTCTGGCTTATTTAACTTAACACAATGACCTCTAGTTCCATCCACACTGTTGTAAATAACAGAATCTCATT
CTTTTTAATGGTTTAATAATACTCCATTTTGTATATGTAACCCGTTTTCTTTATCCATTAATCTGCTAATGGATTGCTA
CCAATTCCTGGCTATTGTGACTAGTGCTACAATAAACACGGGAGTGATAGATATCTCTTTGATATACTGATTTCCCTTCT
TTTGGGTATATACCTAGGAGTGGGATTGCTGGGTCAATGGTAGTGCTATTTTTTAATTTTTTGAGGAACCTCCAACTG
TTCTCCACAGTGGTTGTACTAATTTACATTCCCACCAACAGCATAAAGGGTTCCATTTTCTTCACATCCTCGCCAGCA
TTTGTGTTGCCTGTCTTTTGGATAAAAGCCATTTTAATGAGAGTGAGATGATATCTCATTGTAGTTTTTGATATGCATT
TCTCAATGATGTTGAGCACCTTTTCATATATCTGTTTTCCATTTGTATGTCTTCTTTTGAGAATTTTTTACTCAACTCT
TTCACCCATTTTTTAATTAGATTATTAGATTTAATAGTGTGTTGTGCTCCTTATATGTTCTGGTTATTAATCCCTTA
TCAGATGGATAGTTTGCAATATTTTTCTCCATTCTGTGGGTTTTCTCTTTGCTTTGTTGACTGTTTCCTTTACTGTGC
AGAAGCTTTTTAACTTGATGTGATCTCATTTGTCCACTTCTGTTTTGGTTGCCTGTGCTTGTGGAGCATTACTCAAGAA

Fig. 9.80

Fig. 9.81

TAACATGGCATGTTTGTAGTAAGGAAACCACTGGAAACATTACAGCTACTAGTATATTGGAAATAATCCCTGTACATCTG
GAACAAGAGAAGAATGACTTCGGGTTCTGCTTATGTTTAATTGGTTGTTCTAGCCAGCATTGTAGGTAAAAGAAATAAA
TAAAAGGTAGATAGAAAAGAGAGAAACCTAATTGTTATTAGTAAAGAGACTTCATTGAAAACTATTAGAAAATAATAC
AAAAGTTCAGGAAAATTGCTAGATATAAAGTAAACCACCTTTGTGTAAACCCAGCAAAAGACAATTAGAAAATTCGTGTA
ACTACCTTTGTGTAAACCCAGCAAAAGACAATTAGAAAATTCAGTTTAAAAGCAGATACCATGAATACAGCAAAAATTAG
AAACATGTAGTAATAAATCTAACAAAGGATTTTTAGAAAATCTTTACAAATAAAATTACAAAATTTTAAGGTAAGACACA
AAGAAGATGATAAATAGAAAGATATACTATACTTAAAGCAGAATGACTCAGTATCAGAAAGATTTCAATTCTTTCTGTA
TTCATTTATAAATTCTAATAAGAGCTCACACAAGAGTGGTTTTGTGTTGGTTGTTTTTCAGGAACTTGACAATCTGATCAAT
CTGATTCCTAAAATGTATACAAAGAGCTAAGCGTCAGTCATAACCAACACAATTTTGAAGAAGTCGGGAAGATTAGCCCT
ACCACATGTCAACAGCTACTGTAAAAGTATAGTGATTAAAGACATTGGGATATTGGAGCAGCAATAAATAAATACTAG
GTTTAGAAAATAGACTCATGTGTCCCTAAGGATGTATTACAAATCAGTGTGGTAAGGATACAATAAATGATGCTAGGAGA
TATAATTATTACATGGGGAAATAAAATAAATAAGTACCCTTTCTTACAACATATAAATAATTAAATCCCATGTGGAATGT
AGGTGTAAAGGTGAAAAGAATAGTAAAGATTTTTATGTTTTTTAGAAAGAAAATGTATGAAATATCTTTGTGGCCAGTGGG
CAGGGAAGGATTTCTTAAGACTAAAACCCATTTACGCCTAGTGTTCCATTATTGGAATGCTAAGCATGTGAGAGTTATT
TATTATCCTACTGCTCAAGATCATCGCCAAGGCCTGATTGCAAAAATTCAAAAAATTCGAACCTCAGGCATAAGTGGG
TTAAAAGACAACTATAAAGAAAAATAAATAAGGTTGAATATATTAAAACAAACATTTTTGAATAGCAAATAACACCA
ACAACATTTTTTAAAAAACTTCAGAATAGGAGATATTTATAGTATATAACAACCAACAAAAATTAATATGCATTTTGTATG
AAGGAATCTTTCAAATAAATATTTAAAAATTCAATCCAATGGAAAAATAGGTGAAAGATATGAACAAACAATCGACAGA
AAAAAATGGCTGATAAACACACAAATATTTGCTTTAGCTCACTAGTAATCAAGTAAATGCAAATTAAAAAAAGGAAAAT
TTTATTTTCATACCAACCAGTTTTTAAGAATTAAAGTGTGAAAGTGCCAAATAAGAATAAATTGGTAAACCAATTGGAAA
AGTCAGTTTGACTTTACTTAGTAAAGTTAAACATTTGCATATCTTGAAGGAGGAATTCCTACTTCTAGGCATATATCTAG
AAACATGATTGGATGTGTTTGCCAGGGGACATGCACAAGGATGTTTATTCTAGCATTATTTTGAAATAGCATGATACTC
AAAAATAAATTTCTGTATTTCTATCAACAGAAGAAAGTACAAATAAAAACTGGGATATTTGTGCAATGGACTTACAGCA
TAGTAGATGAAATGTATAAATTCATGAGATACATTAACATGGGTAAATGTTAATAAATTTAATAAACAACATAATAGCA
ACATAAATTTAAGCAAAAACAAAAACAACCTTGCAGAGGGATTAATATTATTTGATGCCATTTCTAGGAAGTTTATAGGAC
ATTCAAAACAATATTTTATGTTTTTTAAAGAATACATCCATAGGCAGTAAAATAATTAAAAGAGGGTGGAAATCCTAAC
ACCAATTGTAGGATTTTTCTGTGAAGGGAGGTAGGGAAGGAGGTAAATGGGCTCAGAGAGGGAAGGATACACTGGGCTT
TCAAATCTAACGGAATTGTTTGTTTTTTTCCCAACATTTTATTTGATAATAATTAAAGATTCATAGGAAGTTGCAAAA
ATTGTACAGATATGTCCTGTGTACCATTACCCAGTTTCCCCCAATGATTATGCCTTGCATTACTATAGTTATTATAGT
CATGAATTTGACATTGATCTCATGTGTTTGTGTAGCTCTATGTCATTTTGTACATGTGTAGATTTGTGTAATCACCAC
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TTTGCCACCATATCTAATCCCTGGTAACCACCTAATATGTTCTTCATCTCTATAAAATATTGTCTTTTTTGAGACTGTTA
TATAAATTAACCATATAGTGTGACTTTTTGAAATTGGCTCTTTTTCTACACAGCATAATGTCTTTGATCCTTGAGATC
CTCCAAGCTGTTGTATGAATCTGTAATCCATTCCCTTTTTATTGCTGAGTAGTATTCATGATATGGATGTAACAGTTTTT
TAACTATTACCTATTGTGGGACATTTTAGTCTTTTCAATTTTTGGCTACTACAAACAACCTGCATATAAGTTTTGCTG
CGATCATAATTTCTTTTTCTCTGGGATAAATGCTTCAGAGTGAATTATGGGGTCATATGGTAAATGCATATTTAGTTTTT
AAAGAAAATTAAAGCTATTTTCCAGAGTTTTTCAAACCTATTTTCCAGAGTTGCTGTACCATTTTATGGTACAGCCATATG
TTGTTGTTCCCTTAATGACATGTGAGAGATCCAGTTCTCTCCATCTTTGTAAGCATTGTTGGTATTGTCACTGTATTTTAT
TTTAGCAGTTCAAAGAGGTGTGTAGTGATGCTTATCATAGTCTTAATTTGCATTTCTCTGATGGATAATGTACTAGTT
TTCTATTGATGCTGTAACAAATAACTGTAAACTTAGTGCTTAAACAACATAAACTTACGATTTTACAGTTCTGTAGG
TCAGAAGTCTAACACAGTTTTTACCAGACTGAAATCAAAGTGTGTTGGTATTCCTTTCTGCAGGTTGTAAGGGAGAATTTG
TTTCTCATCTTGTCTAACTCCTAGGCTCTACTCATATTCCTCGGCTCATGGCTCCCCTTCTCTGTCTTCAAATCCAG
CATTTGTGGGTTTCAGTCTCATATCATATCATTCTTACCTCTTCTGCCTGGGTCTTCCACTTTTCAGAACCCTGTGAT
TAAAGTAGCCCCAACCAATAATCTGAAATAATGGCTGTATTTTAAGGTCAGCTAATTGGCAATCTTAATTCATCTGC
AACCACAGCCCCCTTTGCCATGCCAGAATAACTGGTTCTGAGGATTGGGTGTAGACATTTTTTAGGGAGCCATTGTTCTG
CCTATCATAGGTAGATATGTTGAACATCTTTTCATGTGCTTATTTGCCATCCATATGTCTTCTTCAGTAAAATGTCTAT
GTCTTTTTTGCCATTTTCAAATTGGATTGTTTGTTTTTATTAGTATTGAGTTTTGAGGGTTTTAAATTTTTTTTTGTTGTAGA
TATAAGTCCTCTATAATATTTGTTTTACAAATATTTTCTTTTCAGTCTATAGCTTGTCTTTTCAATCCCTTAGCAGGGGC
CATTCACAAAACAAAAGTTTTTAATTTTGATGAAGTCCAATTCATTGATTTTTAAAAGTAAATTATGCTTTTTGTTGTCA
TTCTGTATAAGAACTCATCACCAGCACTAAGTCCTAACATTTTTCTCCTGTCTTCTAGAAGTTTTATAGTTTTATATT
TAATTCTACCACCCATTTTCAGTAATTTTTGTATAAATAGTGAGGTGCAAGTTGAGATTTATTTTATCTATTTAATTTG
TCTGTGAATGTCTGTATGCATTGCTCCAGCACCACCTTGCTGAAAAGACTATCCTTTCTCTACAGAAGTCTTTTTCCACT
GATGTTAAAAATAAATTGGCTGTATTTGTGTGGGGATGAATTGTTTTATTCTTAATGAAAAGAGGAAATACAATAAAC
TATTAAAGTCTAATAAGGCTAAGGGCATAGTCATGTGTTGTTTTTTATATTTTTCTGTTATATTGAAATATTTTAAAT
CAATAACGTAATAAGGAGGCATTTGCACCCTCAAACCTCTGAAAATATTAAGAAAGTGGGTAATGCTATAGTGAGATGTT
GATCACAGTGGATTGGAGACTAATAGCTGTATGATATTGATAAGGTTACTGGCCTCTCTGGGCTTCAGTTTTATTTTTT
TTTTGTGAAGTGAAGGGCTGAACTAGGAGACCCCTGGAGTCTTTTGACATTTGCATGCTCACATAGTATGTGATGCTT
CTCATTGTTGAGTTTAGAGTAAGAGATGTGGAGCGGGATCTAGGCTGTCTTCTGGTTTACAGCCTAAAGATGAGACAC
TGTTTATATTTCCAGGTAGAGGCTGCCTAATTAACCTACAGGTTTCATAGTGGTTGGGAGCAAGCCCTCTTTTTCAAACCTA
CTTCGGGAGGTTTTATTATACCTGTAATAATCATTATACTATAGAAGATATCATTGTAGTTTCTTATAAATACTCACA
CACTTCTTCTGGATTTTCATAATGGGGTCACTATATGATGCTTCTTGGGATCTGATTTACAATTTATAAATAAGTCTGAA

Fig. 9.82

ATATAAATAAGAAAATACAATGTATAAGAGTCCTAGAGGTTTAGTCAGCAGAATTCCAAATAGGAGTTTAACCCTCTAA
GGAGCACTTACAGTCTTCTTAGAAATAAACACATTGTAAAATGATTATGAGAAAATATTTATCATAACACATTTTAATTG
ATAATTTAGCTGTTATATTTTAATTAGAAAAGAGTTTCTGATTTTCTTTATGGCTATGACTTTAAGCCTGTTATCTAGA
ACACAGTTTATATTTTCTGGTTTCATCATGACAGAAGGCATTTTGAGAAGGGCTAGAGCAAGAAATTAGCAACAGGACG
TCAATTTCTGCTCTTCTTTGTTTACTTCTTCTGGCTAGTAGCAGAATTTTTTTCTATCAGTAATTTTGGCATCAATAAAAT
AATAAAGGATTACAACTTCATCCATCATATGCCAACAAATTTGATAACTTACATGAAATGTACAAATTTCTGGAAAGA
TAGAAACTACCAAAAATGACCCAATAAGAAGTAGAAATTCCAAATAGACCTACAACAAGTAAACAGATTGAATTACTAA
TTTTAAAATTTCCACAAAGAAAGTCCCAGGCCCAAATGGCTTTACTCGTGCATTCTAACAAAAATTTAAATAAGAATG
ACCACAAATCCTTCACAAGTTCCTTCCAGAAAGAGAGAAATTGGAATTTTCATTCATTGTTGGTGGTAATATAAAATAG
CGCAGTCTCTTTGGAAAGCTATTTGGCAGTTTCTGAAAACATTAATTCTAGAGCTGCATATGACCCAGCAGTTTTTGT
CTCCCTAGTATATATCCAAGAGAAATACCATATATCTACAAGAAAATTTTACCCAAATGATCATAGTAGCATTATTTA
TAATAGCAGAAAATAGAAACAACCCAAATGCCTATCAACAAGTGAGGTGACAATCAAAATGTGATATATCCATACGGTG
AGGTATTATTGAGAAATAAAAAGAAATAAAGTATTGATATATGCTACAACATGGATGAACCTTGAAAATATTATGCTTA
GTAAAGGAAGCAAGTCACAAAAGACTACTTCTTGTATGACTCAATTTATATGAAGTGTCCAGAATAGACAAATCTATGA
AAGTTGATTAGTGTCTTCCGACAGCTGGGAGGTTTGGGACAAGAGGGAGTGATTGTTAATGGACCTGTTGTTTTAAATG
TTATTGATTTGATTGTGTTGATGGTTGCATAACTTTGAATATTGTAGAAACGACTGAATTGTACACTTCAAATGGTATG
TGAATTATATCTCAAGAAAATGTTTTTTTAAAAGTAATGTGAATATGTATGTTGAGATAGCATGGTATGTAGAAGTTAT
GCAGGGGAGTGGTTAAGATCATCAACTTGGGTCAGAGTGCCTAAGGTCAACCTCACCATAACAGTTGTGGGAACCTTACA
GCTTTCCTTAATTTCTCTGTGTCTCAGTTTCTTCACCTGTAAGCATAATAATAGTGCCTTTATCATGGTGTCTATCTTAA
CATTAAATGAGCTAATATTTGAAAAGCATTTAAAACAGTATGTGGCACAACATAAGCACTATATAAGTGTGTTGAGAAAT
AAATAAATATAGGAAGACTACAATATTTGTTTTCTTGAAAATTTACCTTTTATTTCAAGAGTTTGATCTTCCTATTCTAT
TATGTTAAAATTTCCCTTTGTAAAAAAGGAAATGTTAGTTACAGTAGATGGCAGTTATTTTGTAAATATCTTTAA
TTTGCTAAAGGAACAGTTGGCACTTCTATATTGATCTCAAAGTAATTTTAAAATATTTTTTGAATTTGTACTTTTCCA
TGAAATCTTAAATTAGAGAACCACAGTTCTATAACTATGTCTTTGTTTCAGTGGCCTTTGGAACCAGATGGCTCAAAAAA
TTAATAACCATTTCAAGTTTTAACATTAGAAGAGTATTACAGATATTCAGCTACCTAGAAAAATACTTATCATATAGGAG
ATGTTAATGAATGTGCATTGGAATGAATGACAAGAATGGATCATAAGTGTGTGACTTGCTCAGTGTGTGTGTGTGTGTG
TGTATGTGTGTGTGTGTGTGACAACGACCTCTTTCTACTGGTATCCTAAAGTCTGTATAGTTATATCTGATATGCTTGA
GTTTGCATGTTTCTGCTAACCTTATTCCCATGAGATTTCACTTTGAGGTATTTATATGGATTCTAATCTTGACGCTAAG
TTTTAGAATCCAATCCTCAAAGTATTAAGGGATAATTGGGATTATTTAATACCCCTCCCAATTTTTCTGTAATGTTTT
CTATTAGTAGTATGGCTGCTGCTATGTTTGCATGTGTTCACTATTTTTATTTCCAATTTCTATTACATTTTTATTTTC
TTCCTAGTTTCTGCAGACTATTTAAATAATTTTATTTGGTCTGTCTCACCCTCTGTGTTGGCAGTTTTAATATCTTTAA
GAGAACTTGAAAATATTTTAGCTCCATAGAGCACGTTTCAGTTTACCTTTTGTATTCCATAGCCTGGATGTAAGGCTT
CTTTATGTTTTGTGAACAGTTAGAACCACACTGACTGCTATGAAAATGCTTAGTAATAATTATGCTTCCTGGATCCTTG
TGAAATAACCTGCTTTACTCAATTTAGTACACAAGTTATTGAGAAACATCTATAGCCTTTTTCATTGCAGTGTGGGACA
GTAGTTCAGTGCATAGGATCTGAGGCGCTGTCACATCTTCTTGATGTCTCAAAAACATAACCATTTGATTTTGCTTCCCA
AGTCTAGATTTACCAACATGTGAGAGACTGATGCTGCAGTGCCAGGAGAATTATTACTGATCATAATCCAAGAAGAACT
GACCAGAGAATTAAAGCATTTAAAGTTGCAAGAAGTAGGTTTATTCTTGACTCAAAAAGGCTTAGTATAACCTACTTG
CACCCCTGTAAGCATCTAAATTATTGTCTTAAGAGAACAGAACCATAATGATAACCACAGTATTGCTTAATATCTGCTG
CTGTTTTTAAAGCATTTCACACAGTATTTCAATTGTGTCTTAATGTTACTAACTTTATATTGCAATTGCTTAGAACTGCT
TTAATTGAAGCAATTTAAATTTACATATCCTACTGAGCCACTTTGTGTGTGAAGCACTAACATGTGATAGTGGTTCTTT
TTTATAGATTACAAAGTTACAAACAGTTGGTTGTAGATCTTATCTGTGAAAATAAAAAGCCTTGAACAAGGCTAATGAG
AATGATTAGAGGGAATTTATTAATAAATTGACTAAGGCGGTCCCCCTCCCTCTATACTATGTCAGCTCACAGCGGGATG
TCTCCAACATTTATTGTCTGTTTAACTCATTTTTTGGCATACAACTATTATGCAACTGTTTCAAATATTAATAGCTCCTC
TCCACAGCTAAACTGTGATCTTGATTTTATAACCAGCCCTAATAGTGTCAAGGACAGAGGTGACAACATCTTAGGAGCTC
AAGACGAAGAATTTGAGGTCATTGTCTGAGAGGTCATGTTTAGTTACATAGCAGGGAATGGTTATATAGGATTACAAC
TGTCTTATTTGAATAATTTACTGCAGCCCTTTGAATGTGTATTTTTTTTCATAGAATTTTCAGTTACTTTGTCTCTTTGA
GATGTTCCCGTGTATTGTTCAAGGCATGGATTTTTATTGGACATACTGGTTGCAAGTGATGATAATCCAACCTTAAATGG
ATCCTATCATTGAAAATTCTAGGGTTAGATCTGGATCCCAGGCTAGAAAAATATCATTAAGATCATTTTTTTTTTCCATT
GCACAGCTCAGTTTTTCTCTTTATTGGCTTCATTCAAGTAGACTTTACTCCTACAACCCTGTCACCAACATAAATAATT
TGCCACAAAGTTTATATCTTATTGTTTCAGCAACCTCAATAGAACAAATAGCTTCTCTATTTGTTTTTAAATAAAAGTCTC
AGGATTGCATCTCACAAGACTGATGCAGGTCATATGACCCCAATTGAACTAATCACTATAGCTAGGACAGTGTCTGCA
CCAAGACTGCCTGAGATGAGAGATGAGAAGTGTGGATTACCAAGGAAGATCATGCTGTCACTATCAGAAAATGGAAGA
ATGAATCTGTTCTCCCCACCCAACAAAAATACATAGTTTTAATTTCTTTCTACTTTATGAAGTATTTCCCAATTATATTT
CTTTTTCTTTTATGCGTGAACACACATTTTCTTTCTCTGTAAACAACTTCTACCTTTAACATTCCCCAGGTCATCACT
GCATTCTCCATGTGAACAGTAAGAATAATGAGAGATCAGCGAAGAAAGCAAACCTTAACCCCTGTTCTGCCAGAACTA
AGGGTTTCAACTGACATCTTCTAGAGTTTATGCCAAAGCCTACCTATTTCTGATGGATGATGTTTAAATTTCCATTGCA
TCCTAAGCTCCACTTCACTTTAGGCTGCCAAATGATTTTAGAAAATCCTCAATGAAGATTGTGTAAACAAAATAGTGG
TTGAGACCTTTACTTATGTATCATATTTCAATTTAGTCTAGCACTTTTCTCTATCATTTGACTCAAAGTTCTCTTTAGCC
AAACATCATTTGTGTCAAAGATCATGGAATACTTTTGTAATATTGAAAGAAGGCCAGCTTCTCCTTTCTTAAAGTGTT
ATGTCAACCTCAGTTAAAATCAGTAGTCCACAGTTGTTTTTGCCAAAACAAGAAATGAATTACATTGGTGGTGAAAGTCC
CTCTTCTAGTGGCAGAGAGAGAATCTAGAATAATTCACCCACTCGGGATATGGGTGGTTTGCAAGGAATGAAAAATCCA

Fig. 9.83

Fig. 9.84

GCAATTCAGTTTTATCAGTGATCCCAGATTTAATCATCTCACAGTCTCTCAGATGATTGAACAAATGAACCTAATAGTG
ACTAATGTAAGAGACAGAAGCTCCATTTATAAAAGTGACAGGAAAATATTTTGAAACAATTACTGAGGTGATTTTTTTT
TTTCATGCAATCCCTTTTACCTGTAGTTCTGTTTGCCTCTTCAATCTAGATCTCAGTCTCTGTGCCTCAGAGAAGGGAG
ATTTTATTAGACTAAAATCTAATGGAGGAAAAAACGTTTATTAATGGTCTTTGAGTTTGAAATTTCTAAAGCCCTGGTA
TAAATACGAGGCAAGTAAGAAAATTCATTCAGTCAAGCCCTGTTTGCTCTGCCTGTTGACTAATGATGAAGATCCTCTT
GACATGGGTAGAACAAGGAAGTGGGAAGATGAAAATGAAGTAAATGCAGATGTTGGTCCCTGAAAAGTTCTCTGCCC
TGCACCCCACTGCAAAACCGAGATCCCTGAGGAGGCAGCGATAATAGTCTGAAATGAATTTGAGGCTCCAAGAGCAGAA
GGACCTGTCCCTTGCTTCCCAGGTGGTACCTGGGAAAATACCCAGATGCTGGAATTTTGTGCATATCCTGGGATAGTATG
ACACCAGTTGAATAGGGGTGATGCTATAATACAGCATGATGTGGTATCTTGGGCCAAGAAAGCCTAAGTTAACTTCACT
GAGTTTCTTATGATCCCTAGAGTGGCTTGCTCTTTCCCTCTCCCCGCTCCCCCAGATTAAAAATGGCCACGAATTCTT
TCTACCTCCCATGAAGGGACAGAGTCTAATTCCTCTCCCTTGAAATGTGGGCTAGCCTGGTGGCTTAATTGATGAATAG
AATGCAGCATAAATGAGCATCTGTGACTTCCAAGGCTAGATAACAAGAAGTTTTGCAGCTTTCCTTAGATTTCTTAGA
ATGCTCCCCACAAGAAAGCCAGCTGTCACATAACAAGTCTGACCATGCTGCTGGAGAGCCTACCTGGGGAGGTGCTGA
GACTGCCTGTAGGGGAGAGGGAGGAGCTCAGTTAAGTCCTGTCTTCCATCCATCTCACCAAGATGACTGGCATGTGAAT
GAAGCCATCTTGGATTTCCAAACCAGTTCAGCCACAGTGAAACACCACCAAGTGACCCAAGGGAAGCAGAATCACACAA
CCCAGTTCTTAAAAATATGACCCGTGGGGCATGAACATTATAAAAAATAATTATTGTTGTGGCCATTAAGTTTTGGATGG
TTTACTTCCCCACAATAGATAACCAGGTCATAGCTCTTCATGTGCTCTTAGAAGAGGGCTGCAATAGACAGAGTTGTCT
AACTTCAACCTCAAGAAGTCTTTTGGTCAGAGAAGAACTCACAAACTGACTCCCAAGGGAACACTTGGGAACAATGCTA
TAGATTATATTTTGTAAATTATGCAATATCTACCTTAAAAGGACCTCAGGGGAGCCTCTTTTTTTAAGTTTGAACAGGAA
AACTCATTCAATTTGCACATCAGCAGTGGGCATTAGCATGGACAAATGTTGACAGAGAAGCAAACAATACTGCCATGAFT
GGTATCATTGGAGTTGTACAACACTGCAGCCTTGCTAGCCACCCACTTACTCTCTAAGTCTTAGACACTACCCAGGAAA
AAGGGGGGATGGTGAGGAACCTTCATTTCACTGGGAATATCCTCATATGGCAGAAGAACTGTGATTTGTCTGAAGACT
ATCATCTTGAACATAAGTTGGTTATTTGCCAGTAGGAAAAAATGGAGTTTGAGAATTAAGAATAGTAATAGAAAAATGAA
ACAGTTATTTTTTAGCACAATTGAGTTTGCAGCTTGTGAGATCAATATCTACCATAGTAATTAAATATGTGAAGTACAG
CCGGTAAGCCAATTAGAAAATCCCCATTATCTGATTAGCTCATCTCTTCTGTGTCCATTTATTCTTTTACAGTGGGAA
TGGTTTTTTCGGTTGTGATCAGAACCATTGTGTCTGCTCCAGTATGFACTCATGCTAGGAGGATAAATTAGTCATCAGGC
CATTCAAGTCATCAGAAGTTCTGAACAAAGGTAATAGGAGAAGCTCTGTCTTGTGGGTGATCATGAGCCTGATGAACCAC
TTCTGCCTTACAATAGAGAACAGCTCTTTGGTGCTTAGAATTCTTGTGGTGTCCCATGCCCTCATTCAAGTATTGTTCTC
TTCCATACGTGCTAAATAAATTATAGCAGATTCTGGCTTCCTGGTTACCATTGATTTCTTCTAATGTGACCATTCCAC
AAGTAATGACAACTTAATCGATAGGTATGTTTATACTGTTAATTAGGTATACTCAAAGCTTGTTTGGGACTTTGTAAT
ATGGCGTCTTTCAAGGATCTCCTCAGAGATCAGATTTCAATTTGTTATATGTTGATTTTCTTAATTTAACTCAGAGTCCT
GAGTATCTCACTGTTGCCTCCATGTACTAAGGGTCACTCTATTCCCACCCTGTCTGCAAGACATTCAAGTGCAGTGCCA
ACCACACTTAACAAGAAGGATCCCAGTAAATAAGAAAGAGGAAGCATTACATATTTTTTGCTTAGGAAATGAAAAGAAG
GAAAACTTTTAACATCTCATTATAAATATCTACCACCCAGTAATTGAGTTATAGCTTATATTGTGCAATTTACAATTCA
TATCTTTAAAGAGCTCACAGTTACCTCTATTTGTAAGTTTTGACAGCAAAAACCTGTTTCATTTCTTCATCAAAACATTC
ATTAATCCCTAATTAGTACCAAGTTCCATGGTTAATGCTTATTAACAGTCCTGTGAAAGGAATATTAATAATATATAAA
CCACAAGGCCCATATAAATCATATGAAATGTTTACTTTTATTAGTCATCAGAGAGATATAAATTTAAACCTCATTTGAGA
TACCCTATACATTTATGAGAATGGCTAAAATGAAAAAGACTAACTATGCCATGTATTAGCAAAGATAGATATGGAGCA
ATTATAACTCTGGTAATGGAAATGGCATAACCAACTCGGAAAACGTTCAAGTATCCCTTAAAGAGTTGAATGTACATCTA
CCCTATGATCTATAAATTCTATTCTTAAATTGTATTAGTCATCTGGATAACAAATTAGCACAAATGTTAATGGCTTATAA
CAAGAATAAACACGTATCATTTCAATTTCTGGGGAATCAGTAATTCAGATGAGGGTTTAGGTGGTTTGGACTCAGAGTC
TTTCATGAGATTTCAAGTAATATGTTGGCCAGGTCTCCAGTCCCCTGAAAGCTTGACTGGGGCTGGAGGCTCTGCTTCC
AATACAGGTTACTCATATGACTAACAAATTGAGGATGACAGTTGGCCAGAGACCTCAGTTCCCTTGCCATGTGGATCCCA
TATAGTGCTGCGTGAGTGTCTCCTGGAAGCTGACGTGCCCTGGAGTGAATGATTCATAGAGAGCAAGGTGGAAGCTT
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TATGATGTATGTATGTGCGGGGAGAAGGGTGTGGCTATGCAAGAGCATTGAGATAAGGAAGCAAGGACTATTGCATGCT
GTTTTTGGAGTCTGGCTACCACACAGGTATTTATCCCAGGGAAATGGAATGAAAGTCCCTCAAACCTTGTACAAGAAT
GTTTCATAGCAGCTTTATTTATAATAGCCAACTAGGTAACAGCCCGGATGTTTACAGGAGTCTGGATAAACAACTGTA
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AAATTAATACAGAAGGGGAAAATATTAAGAGGAGAACTAAGGTACATATATTTTTCAAATAAGTAACACATGGTAATT
GTAGATAATTTTAATAATATCTTATTTTTATATTTATAATCTCCACTTCTGGAATAATTGTTGATTTATTTTATTTTGA
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TTAAGCATCACTTTCAATTGTTATATTCATTGTAAGACTATTTATAATTTATGTAACCGTTCTGTTACTGATAGACCT
TTAGGTGTTTCTCATTTATTTAAATAATGCAGTGGAGTTGTTTATAAAGGGTGATCTTTAGTGTGAGGATTATTTCT
TGGGCTAGGGTTTCCAAAACGTTCTAGATCAAAGAATATGTTTCAATTTTCATATTGCTTTCTACAGGGTTTAAATTA
TTATACACATTTACCAGTAGTCTATTAAAGGACTTATTTTAATAGGTAAGTGAAGTAGGTATTTCACTATGCTTTGATA

Fig. 9.85

AGCATAGTGAACACTTTGTTTCATGTGTTTATTAACCATTTGTAATTGAGAAAAATGGCTGAATTGAGCCTTAAAGAATGA
ATAGCATTTTAAATATCAATTGAAGGCAAAGATGGGGAAGGTTTTTCTGCAATGTTTCATGGCATGGCAAACAAAGTCA
TTCCATGTCAACCTCATGGAATTGTAAGTCATTTAGTATAAAAGCTTAGAGCAGTTGTTCTCAGTCTTGGCTGCACAAT
AGGATCACCTGAGGGAGAATGTAAAAATCTCCAATATCTAGGCCTTACTAGACCAATAAAATCAGGATAGTTGGGTGAA
GAACTCAGGCATCAGTAGTTTCTAAAAGTCCCCAGGTGATTACAATCTGTAAGCAAAATGTGAAACCAGTGGTATAGAG
TGAGGTGAAATGGGATAGGTTGCCATTAGAGATGAAACAGAAGAGATGAAGACCATATGTAGCAAGACACAATGCTCTA
GGATAAATFGTATACTTCCTGGGCAAATTAGCAAATGCTGTTAGGTTAGTGTCCCTTTCTTATCATCTTGGAGTCCCTT
ACCTTTTTTCATGCACTCATGCAGTGTGCATTACTCCCAACCAACAACCTAGCACCTGATCTTGTGGACAGGCTGCAAGCA
AACC GCCCCCCACTGCCACCCACAAGCCTGGAGACCTGGGAGTTTTTTCTCCACACCCCCAGGACATAGGCCTGAAGCAA
TGATTGCTGATGATATGATATAAGTAATCCAGCTTCCTCTTTCTTAATCAAGATAATTCTGACATGTGAACTATACTAG
TCCATTCAAGTAGCCAATATGCAAACCTTTGCCCTTGTTATCTAGGACAAGATGAGCTTGTGCCAGAACATAAATCTGTAT
GCTGCTTCCTTCACAAAGAAATGTTTTCTATGAAAGAAAGGTCAGCTGAAGTAAATAGTGGACTTACTGACTTAGCTGCA
AGCATCTTATTTTCATCATGGACCAGTGACACAGACCAAGGATCTGCAAACCACACTTTGCGTGCTCTGCAGCAGTGAAC
TCATTTTTTAAATATCCTTTTCTTGCCATATCTAAGGTCCCGGCCTTAGTTTTAGATGCCCATTAGTTTTGAGGTTTAGA
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TAAACTGAAATCTCTTTGTGAGATATGTGTTATTTACTGTAGACTTCATAGCTCATAGACTGATTCTAAATTATATTT
GTAAGGTAACCTTAGGTGTATCCACNTATCTGACGTGGAGCAAGTATTGAAAACCTTCTGGTGCTCAGTTTTCTCACTT
CTTGAATGAGGAAAATGAACTACATTATTCCTAAGGTCTCTTTCAGTTCTAAAATGTTATGTTCAACAAGAGCTACTAG
ATTATGTCAGTGAAACAAACATAAAACAGATTATTCAGTAGCAATTGTCACCTTCTCAAACAAATTGACAAAAGTCAT
ACATCTATTGTAACAACAATTCTCCAGGAAGCTAACCTTTGCCCTGTTTTTCATAAGGATATGTTTACTGCTTTTAACTT
TGCTTTGGAAATGGGTAAACCATCTAATTAGGTTGATAAGGTCAACATGCAGGAGCTTTGGAAAGAGATTGGATAATATT
TGTGAGTGTGTATATTTGCCCTAGACATGCTTTTTGTCAAGTTTATTGTACTTTAATGATTAGCTAGAAATAATAAGGCT
GCTGTATATGCTATTTTATTAATGCTGTGTCTATGATGGGTTTTTCGAAGGTAATTTGAAAATACAGAGAAAATATAAT
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TGTCACATCTTGTCCCCAAACAACCCTAAAATAGTTCAATATTTTTTTAGACTCTTTTGTGCTCAACTTATCTAGGAAG
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CTTTTTCATATGCTTACTGGCCTTCTGGGAACCTGTTATATGAAATACTATTCAAATTTTTTTAGCTATTTTCTTTTAAAT
GTCTTTTTTTTGTTTTTGATTTGTTCTTTATGTATTCTGATAAATATGTAAAAAATCAAATTAACAAGTATCTGATTCC
AGTCTGTGCTCTGCTAGTTTTTTTATTAATAATTTATTTTGTGATGAACATAAGTTATTAATTTTAAAGGAAGTCTAATTTAT
CACTATTTTCTGTGATTGTTTCATGCTTTTTTGCGTTGTGTTTTACTATGTGATTGAAAATCCAGTCACCACAAAATGCA
TTCAAATTGACAGATGCTAGGTTTTGTTTCTTTCCCCATTAAAAAATGTAATATTCACATACATAAAATTTACCATCT
TGGCCATTTTTTAAAGTATAAAGTTTTAGTGTTAATAAAAACATTCATAATGTTGTAAAACCATATCATTACCCATCTCCA
GAACTCTTTTTTATTTTATAAAACCAAAACTCTGTATTTCATTAAACAACAACCTCCCCATTCCCTCCTTGTCCCCATTGTCT
GGCAACCCCTGCCAATTTTCTGTCCCTATGATTTTGAATCTAAGTATGTAATATAAGAGGAATTATACAATATTT
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GTCCCATGTGCATTGAGAAGAATGTATATGATGTTATTGTTGGGTAGTGTTCATACCTGTCTATTAGATCAAATTGGT
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TCCAATATTATTATAGAAGTGTCTATTTTTCTTCTCAGTTCTGTGAGTTTTTGCTTCATATTTTTTATTGTCCTATTGG
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TTGTTGGATTCTGTTTTTTAATCTATGCTGCCAATTTGTCATTTGATTGGAAAATTTAATCCATTTACATTTAAAGTAT
GTAATCACAGGTTGAAATAAAGAAGGAAGCAATTTCTCAAAGAAGTAGGTATGCCACTACCAAGAGAAGGGAAATAAG
GTGCTGGGCAGACACAAATAATAGATGTCTATTACAAAGGAAATAAAAATGCTTGACTCATGTGAGCTGGTTGTTGAAA
GTCTCTGGCTCTCAGAACTGGGGCTCTATAGAGGATCACTGCAGGCCTAAGCAAAATAGAAATAACAAGATGCAATTCCA
GACTCTTACTTGGAAGTCACTTTTCATGGGGTTTTAAGGCCAAAGGTGGATCCCTATATTACATCCATCAGAATAAAGCG
CTATCTGATTTTCTCTTACTGGTTCCAAGGAGATGCTCCTGTTGGTAAATAGGGGCCCCAGAATCTCCCAAATTATCA
CCCTTAAAGATCTTGTGTTAGTAATTCATGTTTAATTCCTTTCTTCTTCTAAGTTTCTCCAACATGAGTATAAAATTC

Fig. 9.86

TTATGGCTACAACAGCATTTAGATAATTGATAACTGTGACAGATAAACAAACATAGTTGCTTTTGAATAATTGCTACTTC
TAGTATTAACCTCTGTAGCAATAAGCACTTCTTAGGGAACCTATTCAATTCTTATTTCTATCTTATTTATATTGGGATA
TGTGATTCTCAAACCTTAACATGTGTACAAATCATCTGGACGGCTGGTTAAGATATAGATTGCTGGGCCAAATCTCTCA
GAGTTTATGATAACATAGGTCTGGGATAGGGCCCCAACATTTAACAAGCTCCTAGTGATGCTGCTGTTGCTTGTCCAGG
GATCACACTTTAAGAAGTAGATAAAAAGATACGTCCCTAACATATTAACATGAATAAGTCCTTTATACATCATATTTTG
TCTCATTAGGGAACTTTCAATTAAAAGAACTGTATTTTCTTTGGAAAATAAAGAGCCTTTATTATAAAAAATGAAAAT
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TCCCATTTGCAAGTAGCCTTAACCTCTGCTCTACAGTTTTTTTTATCTGTTTACGCCTAATCAGATTTTGTGGTTGGTTGAT
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GGCTCAAGGACTTCTGGGGCACAGAGATTCACCTTTAGAAGCTTTCTTTCTTCTTGAACACTTTAGGTTTAAATGTCT
CCTTTTAGGACAACTCCTGTTTGGAGTATGGTATTTCTCTATTCTAAGTAAGTATTAAGAATCAATAAGGGCCTCTG
TCCTCTGTTTCAGTTTCTGAAGGACCATGCACCATGGTTAAAATACTTGTCCATAATATGTCATGATAGTGTGGTATGCC
TGCTAACATCACTATTTTCTACTTTTGCAGTTTCTAATTCTGCCCTAACAAAGTAATCTTTAATTCTTCTTCAAGTATTAA
GATAAATAATATTATTAAATTGGTATCTAGTAGTTATACTAAATTCATTTTTTATTAAGTTTCATTAACAGTGCCTGTGTT
CACCACCTTACTCGTAGGTAACAAACATGGGGCATCTTCATTTAAATCAAGTTTAAAAAATATTCTATAAGTTATGTAA
ACACACAACTAGAAAGTATTGAAGATAACACAAGAGAAAGATTGTCAAAGTATCCTGTCTCCTTTCTATGAATTTGCT
GGAGAAAATAGAAAAGTGGACTAGGTCATGAGCAGAATTGAGTTTGTGTGTATTTTCAGGACACCGTTGACGTTTCTTGT
GCTTTAAATGATTTAGAATGGTGCCTGGTGGTAGAAAATTTCTGGAGGCTACATGAGAAGTGATACTAATAAAGTTTT
CATATGTATGTACTCTCTCTCAAAGAGAAAACATTTTATTTGTGATCTTAGCATCTCTCTCAAGGAGATGATATGGAA
GTTGAGCGTCTGCAATGGTTCGTAAATTGAGGGAAGGTTTAATTCACCTTGTTTCATTTAAGTATCATGTACTATGAT
AGGCTCAGGGGAAACCAAGAAAAATTCACCTCCCATGACCTCAATTAATTTCAATTATAGTGAAAGAAACATTAACA
AAATTATAATACATTAAGATGATTATTCTAAAAGCATTTCAGCATAAAGTCCTATGGTGCCTACAGACACCTAATCTCGC
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GCCAGCAACAATAACCAAGCAATAGGAAAATAGCCCCCTGCTTCGCTTCTGTCTGCCAGCTGTCTCACTAGAGCACATG
AACTGGTGGAAATCTAATATGATTTTGAAGTCTTATCGCAAAGTAGTCTGATAATTGTAGTTTTTGCCTTTCCAGACT
CCACAATACAGGTGAGCTCTCTGGAAGAAGGGAGAAATTAATGTTGAGGACCAATTTCATCATATATACTAGACAAAAGA
TGGATGACATGGATAAGGAAATCTATAGAGAAGAGAAGCACCTGAGAACATGAACATTCAAGGGACAATTGAGGAGACT
AAGGATGAATGGTCACGGGGTCAAGGGAAGAACGCAGGAAAATGTAGCATTATGGCAACCAAGAAGAACAAGTTTA
GGAAGAAGGAAGTGGATAATAATGGGAAAATGCTGTAGTGCAGCAGTTATCAATTTTTGTAGTATGACATCTTTTCAGT
AACAAAAATTTCTGTGTACCCCTAGTATTAGGCAAGACTTTGTTACAGAAAATCAACTTAAAGTCATTTATGCAAAAAAT
GAATTTATTGACACATACACTTTAAGTGCAGTGGGGACAGATCTAGCATTAGCTGCTTAAAGTGGTGCATTCGGGTAA
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TTCTAGTATCCCCAGGTCAAATCTTAGGGAAGAGATTGATTGGCCTGGCTTAGGTGAGGTGCCTTTCACTGCACCAATC
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GTCAGTCACACCCAAAACATGGAAAGGATTTTTCCATGAGAATCACAGAAAGTTATGATTTTTCTATTATAAAAAGAGA
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TGTTACTTTGAGAAAAAATATATATATAATGACAAAGAGCTACTATGAATTTAGTGACACTTAAATAAATTTCTAAA
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AATCATGGATTCAAAAAGTAGCCTATGCACAATTTTCAGGTGTTCTTACATAAAGACATGCTCAGGAAATATTTTAAGAC
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TGAGAAAAAATATATAAAAAATGTTATTTCATATAATATAAATAAAAAATATACAAATAAAACGCGTAATAAACACAATATA
TAAACATATATACAAATAAACATTTGTATATATACATTATACATATATATCTATATATATATTTGTATATACATATAT
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GAAATGGGATCTGGACCCGTCCCCAGTGCCCTGTGAAGGGTTGCTCTGTACATTAGCTCCCAATCTGCAAAATGGATG
TTGCTTGACCCGTTTTCTTTCCGCCTCCTGGGAACTGGTTAAAGTCCACACAACTCCTTCAACCTCCAGGGACAGAAA
TGGCATCTACGAGCTAATGTGAGGGTCCCTGGGTGCAGAGAAATGTATTAGGAGCTGTGTTTCCGGAATGACGGTTTCG

Fig. 9.87

TTCTTGGTCTACTGAGCGTCACCGAGCTCGGAGGACACCTCTGTCAATTCAGCAGAGTGACAGGGCTCAATGATTCCAG
CGCCCAGGTGGAGAGAAAGCGGCTGAGGGGCTTGGCCGGCTCCTGTGCGGCCCCCTCAGTCCGCCCCAGTCCCATGCAGG
CTCAGCGCCAGGCTCTCCGCCCCCGCCAGCTGTGGCTGTGCCGCTTCTCAGCTTGGGCCATTTGCAGTGGCCTGGCCTG
CCATAAAAGAGCATCCACCATTAGGATGGAAACGCAGCTGTTAGAAATAGAAGCACCAAAGTGAAACTGTTCTTGGTCC
CTGAGGCCACCTGGTCATCTTCCCCTACATTGTAGCCTACCAGAACTACCTGGGCTGTAATCTAATATTTCTGAAAAAA
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TAAACAGTTAATTAACATGGTTTCTTAGACATCAAAGAGCTTCCAGGTTAAAAAAATTGGGGAGGGGGGTGTATCCTT
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GAATAATTTAATTCGCTTCTTGGTGGAAATGCAGCTTTATTTTCTTGTTTTAAGGATTGAATGCATCTATGTATTGATT
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CACAAATAGCAGTTTTACTAAAGCTTTGCTTTTATTTTGAGTTAATATATTAGATTTAGTACATTTGTAAAGTAGTTAA
GGAACTGGCAGTATTATTTTTTTTCCCAATTTAGAATCTACATTGACATTTGTAGATTCTAAGTTAGTATGGATCTCAA
AATTTGTGCTTTCAATTCACATTGAGGGAAATAAGACTGAGCTGCTGTGTATTTTCTTTCTTTTCAATGTATTTTGCC
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AAAAAACACAACCATTTTTATACTTGCCTTCATGATCTTTCTGTCTCCCTCCTTCCCTCTCCCTCCTTTTCTCCCCCTCC
CTCTTTCTCCCTCTTTTCTCTCCTCCCACTCTCCTCCCTCCGGCTCTCTTTCTCATCTCTCTGCCAGCGTAAGTGTA
TAACTTCTTAACCTTATCTCAGAAGATGAATACTATGCCCAAACCTAGGAAAAAAATCCCACTAGAAATGATGTTGCAA
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CCATACAACAATAGGCCCTTTGCTGGATTGAAGTGTTGCTAAAATTCTAGTGTGTTTGTTTTTTAGATGAATGGCAGAA
TTGCCAACAGGAAGCTTTTCCAAAATCTCCATTGTTATCCAGAGAGAAAAGAATGACATAGCAATGTTTTTAACATTTT
AACAGCATAATTTTCTTTGTCTCTGAAAACGGGAAGCTACGTATCATTAGGAATTTCTGATTAATTTCTTAAAGTAT
AAGGGTTCTAGTGCTATACAGGGAGCAACCCAGGCGACCTGCTCATTAACACTAAAAAAAAGAAA
AAAAGAAAAAATACTTTCTTGGGGACAAGCAGGATTTCTAAGGGCTTGCTGGCAATGTTATTTGACTGTACATAGAGG
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CAAAGATCTCAGAAATCTGGTTTAAATTTATCAGCTTTGAGTTGCTTTTTCTTTCACCTTTTTCATGCTTGTCACTGGCT
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CAAACTTACAAGACCGTAACCTACACAGTTCAACCCTACATACACCCTGTGATATCTCTGAACTCTTCTCTACTCCTCT
CTTCCCTGCTCACTTGGCCCCAGCTGTGCTTCAGACACACCAAGGTACTTCCAGACACTTACAGGCTTCACTCCTGCCC
CTCCATTAAGCTGTTACTCAAATGTCGTCTTTTCAAAGAATCTCTTTCTGGGCATGCTATCTAAAATTGCAATCCCTAC
TGCTAATACACTCTATTTTTGTCTTTGCACATATCACCATCTCACAGAATGTATTTTACTCACTAATCACTTACACATT
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CCCATTTAAGTTTGGGTCAAATTCAGTATACTGACTTATTAATGAAGTCATTTGGAATGAGTAAGAGTCCTAGGTCTA
TGTCGATCACACTGAAATAAGAGCAGACCATTGATTGATTCAATACTAAATGTGATTGTTCTATAACTTTCTGAACACT
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ATTCATCCTCTGTTAGCTGAATATTCGTCTTAAAGTAGGGCCCTGCCATGCCATTTCTCCTGCTAAGAGTCTGCATTGGT
TTCTACTGGATCAAATCCAAGCTAAGGCACTCAAGTGCCAAGCAAGGCACTCAAAGCCTCATATAGTCTCATTTCCTCC
TTAACCATCTTAGGTTATTTACCTCAGCTGCCTGTAACAAATCCTCTGCTCTAGTCAAGAAGTTTCTCTTCTTTCTTTC
TTCTTCTCCTTATTTAGCCACCCCTCCTGCGTTAGGCACCCGTTTACTAGTCTGCCCTGCTTCAGGGACTTCTTATTT
CTTCTCTGATGTTCTTTTCAAGCATGAATGTCATTCTTTTTTTCTGCTTATCCATACACCTTATGGACCTATCTGTAA
GATCCACCTGTAAATTACACACTTTGTTTTTCATGTCCCTAGCCTACAACAAATAATCTCTCTCATTTTTTCTTTTCTTAA
TTACTTTTGAACCTTATCAGGTGTACCACACAATTTATTAATCCTTATTTAATTACAGACTGTAGCATTCTATAATTGTT
TCTAGTGTAGCCTTGTCTCTTAAACAAGATTGTAAGTGCTTAAAGAGATTTATGTCTTATACTTTCCCGCCTATACTT
TGTAAGTAGAGGTTGTTTTTATTAAGAGATGCCCTACAAATACTTTGGCTTTTTCATCCTGTATAGGTATGAGTCCTGCTA
TTAATAGAAACAATGACATCATTTTATGATAGATAGTAAATGTGGTTTATCCAGCAGCAGATCCTTCCAATCTGATAT
TAGTTTTATGCATTTTATGAGCACACATATTTTCAGAAAGTCACTGGAAGATTTTGCTTCTGTTATATTTGAAATTTGA
ACACCAAACCTTCCACATTGTGAAATGTTTGCATGAACACTTTTGAGAACTTTAGATGAAAGGTGTAGTATAAGTACAAA
GTATGCATCTTCAAAAAGCAAATGAAAATGCAAATATTTAGAAATTTCAAAACAAAGCATGGGAAATTTTGGGTATAT
TGCAAGGCCAAATAATTCATCATTTCCATTTCTAGAGCACTAGAAAAGGTTGGGAAATCTGTCCTTTGAAGCCTTAGAGT

Fig. 9.88

ATGTATATTTTTCTCTTTAGCCCTGTGCTGTTTCCTTGAGGATATGCCTGTAGCAATAAAGGTAATCGGGAAGGCTTTG
AATTTCTGAGACAGTGTTAAGCATTTTTTAACATCAGATTAAAGGTGGCAAAAGCTAGGGATGATCTACAGGTGACTTC
AAATCAGGGGTTACATTGCAAGTCCGTCTTTTGGATGAGATAAGTCAAAGTGCCAGTGAATGTGTTTTGGTGGCAAGAA
AAGAAGCTGAGAGGGTCAGGATGCAGAAGTCAGCATTTTTTCAAGAAACACTGGGATAGAATTTCTTTGTGTGGAAGT
TACCATGTCTGGAGATACCTCACTTAATGACATTAGTTGAATACTGTGCAGTTTGTCCAATTTTCAAAATGAAGTACAT
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GCCCAGGTTGGAGTGCAGTGGGGTGATCTCCGCTCACTGCAACCCCCGCTCCCAGGTTTGAACAACCTCTCCTGTCTCA
GCCTCCCGAGTACTACAGGCACACATCACCACGCCTGGCTAATTTTTTTTTTTTTTGTACTTTTAGTAGAGACGGGGTTT
CACCATATTGGTCAGGTTGGTCTCAAACCTCATGACCTCAGGTGATCCACCCGCTTGGCCTCCCAAAGTGCTGAGATTA
CAGGCGTGAGCCACCATGCCCCACCCTATTGCCTTTTCAGGAAAGTTTTTGGAGTGTTCTGAAGGTTGGGGAGGATCCA
CATTCTCTATCTTTAGAAGCTTCTCTTTAGTGTCCTTTGAATGCTGGCTTCAGTTCAGTGGATAATAGGTGGATCAG
GCTGGGTGCATTTTTTAGATTTTGTGGTATCAGAATTTGAAAACAAGATCTGCTCCAAGGGTAGGGGCAGGTCCCACTGG
TAGAGACAAAAGGATGTTTTGCCAGTTTGCAAGCAGAGTGCAATGTACTGAAAGAGGAGTGCTAAGTGCAAAGTTGCAC
ATCTAAGCCAGCTGTCACTGGGCAAAGTCTTGCAAAGTCTCACAGGTGGCACCCCATGGGCCGGCCCATCACAGGAAGA
CAGAGCCTGAGTGTGGCCAGTTTGTGGTGCTCCAGGCAGTGGGTCTGAAGCTGGGCTCAAGTAGGCCTGGCAGCTGAC
AGCGGACCTAGCCTTTGAGCTGGGGATCAGAGTTGCAGGCTTAAAGGAGCTCTCAGGCAGAAACCTGGAAGACAAGGAA
GGTGGGGGAGTATGATAAGGACCAGCTGCTGAAAACGGGGCACACGTGGCTACAAAATAATAATAATCCCCACAGATA
ATAATAATAAAGATAGCTAACACTTATTGATGCTTACTATATGTGAGAAAATGTCCCAGGTCCTTCACACATTTTAACT
CTATCAATCCTTAAAGGCTGGTACTGCTATCATCCCCACCTTATGGGGGAGTAAACTGAGTCCAGTAAGGTGGAATAG
CATTGCAGAGCTACACATCTAATTAGTGGTAGAATAGACATCTTAATCCAAGCATTCTGGGTAAAAAGTCTGTGTGTT
TATCCTCCAGGCTATCCCACCTTCAAGTACTAAAAGGTGATACAAAATAAATTGAGTCTTTACAGTTTATTGGAAAGGG
CAAAAGCTTTGAGCTAGAGAGATAAGGATTTGGATCTCACCTTACCCTTGTATTTCTGACATTTTGGGAAAATCAT
GTGATTATCTTAGATTTGCTATATCATATGTAGAATGCGGGAAATGCCATCCAGCCCATTGATTTTGAAGAACTAAGTG
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AAATGCCCATTCGTATAGAGCTCTTTCAGTTACTTCCGATTGGAAATAGGATTATTCCAGGGCACAAGTCTGGGCTAAG
CTTGAGATAATTAGGGAAGCCACAGGTTTCAACCCTTGTGAAAAGAAGAATAGATAGATAGATAGATAGATGACAGATA
GAGACAGATAGATAGATAGATGACAGAGAGATAGAGAGATATAGATTAGATAGATGACAGAGAGATAGAGAGATAGCTA
GCTAGCTAGATAGATAGATAGACAGACAGATAAAAGACACCACAGGCTATAGTGAGAGGTGAGACAGCACCTGGC
TGTTTCTGCTGCTTACCCTTAAACAATGTGGTGCTTTCTTCTTTTAAATTTACAGAAAAATGTGCTGGCAAGTGCTTT
TGTATACAGTTAATTCAATCAATGAACTTTGCATCTTAAAGATGTAACTAATGCTCACTTCAGGAACAATAATTGGC
AAATTTAAAAATTATTCAATTTTTATAAACATGTTAATCTTTTGCTTTCAAGATTTTTTGTGTTTCACTTTTGTAGT
ACTCAGAACTGACTGAAATGATTCTAAGTTTGAATTTCTATAATTATGCTTGAATTTCAAAGCCTACCTTGCATAGGAT
GGTGGCTAGGGCATAAATATTACATCCTATCTCCAGTTAAGGCATGGATACCTGCACCATCTTTCATATGAGAAGCATC
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TAAAGCTATTACTGTATATACTTTTTCTGTACTTAAAAACATATTTGATAGAAATAGCCACGTGTTTCGCTGTAGAAA
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CATTGCATTTTTTATAGCGATACTAATTTAGGTAATCTGCTATTGTTAAGTATTTAATTTATTCACTAGGCCGGGCATGG
TGGCTCATGCCTGTAATCCCTGCACCTTTGGGAGGTGGAGGCGGGTAGATCACCTGAGATCAGGAGTTCGAGACCAGCCT
GGCCAACATGATGAAACCCGTCTCTACTAAAAATATAAAAATTAGTCGAGCGTAGTGTTGGTGGGTGCCTGTAATCCCAGCT
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GCCAAATACATCAAATATCTAGAAATAGCATGCTGACTATACTCCCTCTGGCAATAAATGAGTGCCTCAAACTGTATT
TTTTTTTCTTTTCATGTACCATCTGATAGACAAAATAGGCATCTCCTTATTTTATTTTTTTATTCAATTTGTTTTCTCGCA
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TACTGGGACATATTTATTTTCTTACTGATTTTTAAGAGTTCTCCATGTTGTACCTCTTTAACTATAAAATATGTAATA
TGTAATATTTTCCAGGTTGTATTGTATATTAATTTTTTTGATATATAGAAGTTTTTGTTTTAAATTTGATTAAATCA
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TGCACCTGTAGTCCCAGCTACTAGGGAGGCTGAGGTGGGAGGATCACTTGAGCCCAGGCAACAGGTTTGATTGCACCAC
TGCACTCCAGTCTGGGTGACAGAGCCAAACCCTGTCTCAAAAACAAATAAATAGATAAAATAAAATTAATGAAATCATTT
TATTCATAGTTTATGCACTTAGAAAAGCATCTGGCATATATAGTGCTCAGAAAGATCTTCCTCACCTTAGAGAAATAA
AACGTTACATATATTTTCTTTGAGTATTTCCATAGCTTCCTTTTTTCATATTTAAAGTTTGAATTCATCTGGACTTTAT
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CCATATGCTGATAAAAAATGTTAACTCTGTCAAATATCAAATTCCTTATGTATAGTTGTATCTCTTTGGTGTCTCTGTT
AGTTCTATTGGATTCTCTTCTAGTCTGGTGCCAGTATCAATTATTATAATTATTACAGAAATTAATCCTTTGTTAATAT
TGTCTTTCAGAAATGTTCTGGATATATTTACATTTATTTATATGAGGTCTTAAATAACACAAGCAATGGGAGACAAGGG
GGAATTGCAATTGGGGGTAAATAAAGCAAGTTTGGAAAAATGTTGATATTGTTGTAGCTGTGTGATGGGAATGTGGGAC
TCATGATCCTATTTACTCTGCTTGCATGTATGTGTGAAAGTGACGATAACAAAACCTTTATATAAATAATTAGGTAAATA

Fig. 9.89

Fig. 9.90

TTTTATTTTCCTCATGATGACTCAGGGAATTTTATCCTGAGGCTCTCAATGGATCTGTTATAGTCAAGCTGATGATGAC
ACATCTCAGAGGCTCATTCCATATTGGGCCCTCAGGCCAGCAAACCTACGCACCATGCTGTCTTATGATTACAGGACGAA
ACTGCTCTAGACACCTCCCACTGCCCAGGCAGGAAGGGCACCATAATGAGGCAGTGGGAGGGTGTATGGCTAGGAAAGT
TGCTAAAAGGAAGCTTTTGTGTAACTTTCTTCTCTGCTGCAGGAGGCTAACACCAAAGCAAAGTATTATCAAGCAACA
GACCCTACATTTATGCAATATTAATGAGAAGGTCCCTGGACTTTTAATTAGGGTGGAGAGTTGTGTTTTAGAGAGCTGA
AATTATTTTAGGTGGGAGGAAATGGAATCCTGGGAGAAAAGTAATTAATTTATAGCTCAATGACTGAACTGTGCCATTT
TGAAATACTGGTCAAGGTGAGCGTTGAAAGAGTGGGTACTCTGGCATTTCATACCCCTTGGGAGTGAAGAATTAGGCTA
TTTGGGGAGCCAAATGATATGCTTGTATTTCCAAAAATGTCATTTATTTTGTAAATAAATTTGTGCTGAAATAAAAAGTG
GCTTGTGTTCTCTATGTAAGTGTGACCGCTGCATATTATTCATTCATGGTTCATTTACAGCATACTTTATAGGTGTGG
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CTTTCATGTTCCCTTTCCCTCTACGTGGGGAAAAACATCAGTATATGAAATGGCATTGTAATAACTTAAAGAGAAGTGTT
CACAAGAGCAGAATAACTCGGAACAGGCTTTGAAGCCATTAGGTGTATGAATCATTTACTGCCTCCTCGGGGGTCCCAC
ACCAGCTATTCTGAGATTCTGTGTCAGAGCACTTAGTGTATTGTGTTATTTGTCTCTGTCCCCCACTAGAATGAGAGTTCC
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TGTTTTCTAACCTCAAGGGACTTAGAGTCCAGGGCAAGGTTGGCAGTGACGTACAATTAAACAGATCATTTTGATGTAA
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GGAATGGGGCTGGGATATTGGGAGAATCAGGAACTCTTCCAGGAGGAGATGACACCTGAGTTGAGTCTTGAAGCAAGA
CATTCCTCAAACAAAGGAAGAAGAGCATGGCTATGGGGTAAAGGAAGAGGGGAAAAAGTCCAAGACAGAGGACAGGGAGG
TGAGAAACCGTATGGCGTATGCAAAGAATTACAAGTGGTTCCTTATAGCAAGTGAGTGAAACAATGCTGTGTGGGAGGT
CAAGGGATGATACTGCAAACAGCAGACAGGGGGACTCCTACTTACCATGTAAAGGAGATTTATTTGGACTTTATGCTGT
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GCCTGAATTTGTGTAGATAAACTCTGGAAGACTGTGGTAGTGGTAGACCCCTCTTAGTGTATGAGATATGTATGTGGC
GGGTTAAAGTATACACATGGCAGCTAAAACAATGAAAAGTGTGGATTATCAAGAAATATAATCTAGAACAGAGGCTCC
CAAGTGCCAATCAAAGTGGTGGTGTCTTTCACCCCTTCATGATTATTATCTGGGGTATTGGTCAAAGTAGATTCTTGA
GTACCTGTCTCAAAGTTCTGATTCATTGGTGGAGAGTCAGCCATCTGCATTTTATGGCATTCCTTAGCTAATTTTGAT
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GAACATAGGGTTATAAGGATAATTTACTTTTTTACAAGAGGAGAGACTCAAGAGAGTATTTAGATCTTGAGGGGAGAGAA
CCAGTTAGAACTAGAGCAGATGGAGGAGGAAAGACAGGCATAGGTAGAAGACAGGACTCCTCATAGGAGGAGAAACACC
TCACTCATCCTCTCAGACAGCGGGAAAGGAGGTAAGGGAGAGTAAGCTGGGTGAGGAAGGAAGCTGAGGCATAATTTTT
GATGGGATGTTTCAGAGAGACATTAAGAGGCTCAAATAGTTGAAAGGGTACAAAGCTGGTTGAAACTAAACCTAGGTAAA
GAGCCCTATTGTTAACTTACTATTTTAAAAATATTTTCTTGAAAATTGTGACAGCTTTTTCTTTTCTTTTCTTTT
TCTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTT
GCTCACTGCAGGCTCCGCCCCCTGGGGGTTACAGCCATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCGC
CCGCCACCTCGCCCCGGCTAATTTTTTGTATTTTATAGTAGAGACGGGGTTTCATCTTGTAGCCAGGATGGTCTCGATCT
CCTGACCTCGTGATCCACCCGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACTGCGCCCCGGCCTGTGAC
AGCTTTTTTCCACCCATATTGCATTTATTTTCTAACTTATAATAATATGAACCTATTTTGTAGATTGAGCTCCTGTTGGT
CCTATAAAACCCAAATCTCTTGCTTGCTGATTCAGTTGTTTATTTGCATTTGTCTCATGTTCCATGTGTTTTGCATTCC
CGGAAACTAGCACAGGCAGGGAGGCCTCAACACCTAATTTATTTCTTAGGTGAAGTGGAACCTGAAGATTTTCTTTT
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AGAGAGAACTTAATTTTAGCCATCTGTTTTCCAACCTGGCTTTTCTCAGTCTATAGATTCCCAAAAACAAAGTCTAACA
TTGGGTACGAAATAGCAAACCTTGAGCACTGAAAAACAAACAAATGAGAAAGCAATCTAACAATTCCACGAACAAAAAAC
TAGCACTTTTGTATGTCCTCTGAAGTAAGAAGAGATGAGTAACTTTTGTATTCACTCAAATACTTTCTTTTAAATCCAAC
TTTTAAGTATTTTTTCTAAGTGTGAAAATTTGAGAAGCCAAGAAACATAGAAATGAGACAGAACTACCCATAATACTG
TACCTAAATATACTACTAATATTTTGCTAACTTTTTCAGTCTTTTCTCTGTGCATAACTTTTTATAGGATAATATTAC
ACATACAACCTTAATCCTGCCTTTTTTCACTTAATATTATAACACAAGCTTTTTCTCATATAATTGAAATATCTTGCAAA
ATATATGATTTTAAATTACATAGTACATCATGAAGGTTCTCATATTTTATTTAGCAATTCCTTTACTATTTGACATCTAA
ATTGTTTCCAGATTTTCATTATTTTAAAGTAGCACTGCTGGAATGTTTCTGCATACAAAGCACTTTCCATATTTCAAGTT
ACTTCCTGAATTATGATATAAAAGTGATATAATACCATTAAAGCTTTTGGTACTTATTATCAAATTCCTATCCAAAGACT
TATGTCATTTGGAAAGTTGATCAGCAATATATGAAATGCATATCTAATTATATACACCATTTTCAAGTATTTTATTT
TTAAAGTAAATAATTTTTAAGCAAGACAAATCTTGAGATTTTAAATTTTCAATTTTATTAATAGTGAGATTGAATATAC
CCCATATTTTGATATAATTTTATTTCCAATTTTGCAAAATTTCTACTTATTAATTAACAACTTCAATTTTGTAGATTTA
TACCTCTAATTTTAAACATATTTTGTAATGGTACAATGTATGTTACATATTAATACTATGGCAAAGGCAAAAGAATGA
TAATGAATTGCTTTCAAACCTTAAATATATATATATTTTTTGAGATGGAGTCTCACTCTGCCACCCAGCCTGGAGTGCAA
TGGTGCAATCTCAGCTCACTGCAACCTCTGCCTCCTGAGTTCAAGCGATTCTCCTGCCTCAGCCTCTTGAGTAGCTGGG

Fig. 9.91

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TTACAGGTGCCTGCCACCATGCCCAGCTAATTTTTTGTATTTTTTTTTTAATGGAGACAGAGTTTCACCATGTTGGCCAGG
CTGGTCTCACACTCCTGACCTCAGGTGATCCACCTGCCTTGGTCTCCCAAAGTGCTGGGATGACAGGCACGAGCCACTG
TGCCTGGTAGAAGTTATATTTTTTAATCATACAACCTTGCAATGGAAGAGACTTTACAATGTACCTTATTTCAACCCTG
TATTATTAGTTTGTTCATACTGCTATAAAGAACTGCTCAAGACTGGGCAATTTATAAAGGTAAGAGGTTTAAATTGGC
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TCACAAGGAGGCAGGAAGGAGAATGAACACAGGAGGAACCTACCAATACTTATAAAACCATCAGATCTCATGAGAACTC
ACCCACTATCATGAGATCAGCATGGGGAAAACAGCCTCCATGATTCCATTACCTCCATTTGGTTTCTCCCTTGTACGT
GGGGATTATGGGGATTATAATTCAAGACGAGATTTTGGGTAGGGACACAGCCAAACCATATCAATGCCCCAGTTGAAGA
AGCAAAGATTAAACCCATAAAACAATTATTGAAAAAAAATGTTAGCTAGTAGATGAGTTTGTGGTACAACATACTTTT
TGTTATTGGACCTCAGAGTCAGTAGAAATCTATGTGACTTGGCGCAGTAGGGGAAGTTGTTATTAGAAAGAAGGGCTTT
TGTTGGACTTTGAATAGGAAGAGTTCTGTAAAATGAAGAGAACAAAAATTAAAGGGCATGAATCTAAGGCCAAGAGGTA
GAAATGGATGTGTCTCAGGTGCGGCAGAGATGAGACCAGCCTGGAGGGTAGAGGGCTGGTGTGGGAGATACAGTGTGA
GGTAAGGCCAAATTAAGCATGCAATCACAGATCACTTGACTCTTCTAGTTGAAACACATCACTCTGGAGTTCCTACTG
AGCATAAGCTACAACCTGATTATATAAAGCCATCCTTGTGGTTACAGGTAGATTGCACTGACTTTGGCAAGTTTCAAGA
TGTTAAACTGTGCTGGGAAGACTACAAGATGGGTGACAGATAAAACTAGTTTTAGGAAAATGCATGCACAGGGCAATA
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AGGCAGACTACTTATGTTCCCTATACTTCAATTACCCTATTTGTACTCTACGTTCTTTCTAAATCTCAGATTCTATAGC
CTCTAGCAGGAGCTCAGATAAAGCATAGCTTCAGATAAACAGCACTCTAGAGAAAAGAGAGAAAATGTAAATAGCAGGA
AATCTCTATTGTCTCTTTAGATCTGGAATAAGAAGTCCTTTCTTTTAATTCACAATTTTAGAAGGTATGAGGAGATAT
CCAGAACTTTCTTCCCTTGTTTCTGCTACCAATGTAGCAGAGGGCCAACCTGAAGCAGATGTTTACATTATCTATCTTAC
GTGTGTAGAGGGCAAATTAGTGGACATTTAAGAGCCCAAGGCAATAATTCAACCATTTCCCATGAAAAATCTGTTATTCC
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TACTTTGGTGAGTAAACCTTGATAAATTAAGCCTGTTTTTATTATATTCTTTGGAGTTGTCTTTGATTGTGATCAAGC
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CCACATTATCTTTGCTGAGAGAGAACTTGCAGGAACTAAGATAACTGCCTTCTGAGAGTCAACCTTTTCTATCAACA
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GGCAGCAAAGCACTATGGAAGAGTAGACAAAAACGTGGATTTTGTAGGCAGACAAATCAGGTGTGAATACTGGTTCTGCC
ATTCCTGGGCAATTTACTTCATTCTGTTGCTGCTGATTTCTCTGTTTGTGCCATGGGAAAACATATTAACAATGTCTACC
TCACAGGAGTATTGGAAAGTTTAAATGATACATTATAGAGATTTTTTTAAGGATAAATGAGGTAGCAGAACATGCAATCC
TCTTCATAGACTGCAGAGCACATCTACCATTTCCTCTCCATCTCCACTCTCCTCCCTGCCCTACCTTGGGGGGCTTTGA
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AAAACCTGGTTAATTCAGCAAAGCTTATCAGGTCAAATCCATTATTTGTCTGATTTGACTGATTTGTTACCATTGAGTC
ACTAGCCCAGTAGGGCAACTATTCCATGGTTGTCCCTAAGGCTACTCATTAAATCCTGGATGAATAATTAAATATTTTG
AATAAGTTTTTCTCTGATAATATATGTTTCTACGGCTGTTATCTAAAGTTTTTTCTCCCTAGATATGGAATATTTTCAT
TCAGTTTGTATTAAATTTCTGTCCAATTTCTAAATTTACATGAGTAACATAATTCTGCATTTTCTGGGACCTATAGGATGC
TAATTTGTAAAGGTGATTCAATTTCTGGAGGTGTACTAGCTGAGAACTTTCCATTGTGGATCAGCTCCTCCCTTCAAAT
CCTACTCCTTTAGAAAAAATCCATACACACTCAGAGAAACAGTATTTATCTTAGCAACTCACATTTGATTGTGCATTTT
TCTTTAATCTTCAGGCAAGCATTTCTATCAACCTTGGAAGAAGGCTTTGTCCCTTGTTTCCCTTTGAGTCCCCAAGTTG

Fig. 9.92

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CATCCAGAGATATTCTTAGCCACAGTGAACCTTACTTCTCTGTTTCTACCTCTACTTTGCTGTTACGGGACCTCTTACC
TCCCGCAAAGTGTCTTCCTTTTATTTTGAGAAGACTTGAGAGGGTGACTCACATATATTCCAAACAAGTATTTTCAGCC
TTTAAAAAGGCTGTGTTCCCTTGCAGGCTTCTCTGCTTTCATTTTGTATGTTTTTTTAAAAAATGATACATAGTTACTTT
GTTTTTTATCTTTTAAATGTATAATTCATACTACTTTGTACTTTAATATTGTCAATCATTTTAGCAAAACCAGCTCCTT
CAGACCTTAATCACTGTTACTCTTTTCCTTAAGTCTCAGACACATGTTTTTTTGAGAAGCTTACAACAAACCCAAATGAT
AGAACTACATGCTGCTGTTAGCATCAGCCTACACCTACACTATTAGCCTAAACCTGCAATATCAGAGTTTTTGTGGTTT
GACCAAGCCTTTTATCCTCTTTCCCTTACAGTAAGTTCTTTCCAGAAAGAAAATATCAGCCCACATGCAAATATATCT
TATAAATATGTAGGTTGTGTCCTGATGTAGCAGAATATCATAAATGAACACAGCACATATAGCTATTTCGATTTGTTCTT
TCTTCAAAACTCAAAATGGAGAATCTCCACTGTAGTGTCTAGGAGAAAGCAAAGATTAGTTAGGTAATGAGGATGTCAA
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TGTCCCAACAAGAATCAACATATACACAATTCAGTTTGCATCTTCATTTTATACATGTAACCTTTAGGTTATGGCTATCA
TATCTGTTTTTTTTTTTTTTTTCAGCCACTAAACTGTAGAGTTGAATATTTAATGGAAAACAGATGGTGCTTGAAATCTC
TAACTTACATTTAACAAGCTGTTTTTGAAATGTTGCTTATGTATACTTTTTTTATTGCTTTAAACACAGCACCGTTTT
AGCCTGTTCCGATGAATTATTTAAGAATTAAGTGTCCCAGTCTAAGACAGCATTTCAAAGTGCAAGTGTTAATCATAAC
TTGATTAAACATTTCTTCTCTTTTCTTTTTCAGCATTCAGTTGGCTTTTGAGTGGATACGTGCAGTGAGATCATTGACA
CTGGAAACACTAGTTCCCATTTTAATTACTTTAAACACACGATGAAAAGAAATACCTGTGATTTGCTTTCTCGGAGCA
AAAGTGTAAGTAACCTTTTGTTCATCTATTTTCTAAACACATGTACATATAACATTTTAGTTTTGGTTTTGGATTTTA
ATGCTATGCTATCATGATTAGGCTTGTGGGAAACGTTTAGTCAACTTTTCAAGTTCTCTGACTGTACACAGCTTATTAACA
AACAAAGACTAGTTTGTCTGTGTGACCTGTTAGCATTTGATGAGAATTCTTTCAGCTAGTTTATATCTTACCATTCAAT
CATATACGTCTACAAGCCTGATGTTGAACACTCTAATTTTTGAGTGCATGCAGAAAATATATGATTAAAGTCAAATA
TAGGGAATTTATGTCTTGAATATGAAACCTTCTGGGCAAGTCTAAAAGCATAAATTATACCTTATGTTATTAGTTACTTC
AACCAGTCACATTTCTGAAAGTTCTTCCCTTCTATGAGCTTTCTACCCTGGCATATATCTCTAATTTCTTCTTTTAATCT
TTTTTTAAAAACATTTTAAATTGAACATCCTCAGGGCTCTACTGAAGGTTAAACCTTATTTCCAATTATGTAGTGTCT
TAAGGTACCACAGTCATGATACTTTTTTAAATTTATATTAGTGTATGGAAAAAGCAAATAGAGCAACTCAAAGAAAGCC
ACACAAGGAAAGGATGAAAGCAGACTATTAATTTCCATCAAGCAAGCGCTAAGAACACATCCCTTTGTTTCATTTACCCA
AGGGGGATAAGGCATCCAAACAGATGTACTTGTGACGTAGGAACATTAATTTGAAGGCATCAGAAAACCAAAATGCA
AACACATTTCTCTAGTATGGGAACACTTTGTGTTATAACCAGTGATTCTATTGTGGTGGCCCAAGGCCATCTTTCATTC
TTTCTATGGAGTTACCATCCAGCTTTTAAGGGTGAGGCATGAGTATGTGCAGATAAAGTATAGTATGCCCAACTGTGTT
TGTTAAAAAATGAGGATATGCTGAGAGATATGCCAATCCTTTGACTGAATACCGTGTTTAATAACCAACCAAGTAAAT
CAACAACTCTGAGGAATTGTTCAAATTTATGTGTTTTACTGGTTGGTGTGTTGGTTGTATGGATTATTGCATAAAAGAA
GGGGCCATGGATTTGAGCCTGGAATTCCTTTTATCATCATACAAAGTCACTCACTGTAATAGGAGGTGACCTGCATACA
AACTACCAAACCTGCAAACACATCTTTCTCTTACTGAGTTTTCTTATTATAAATTAAATATGAAAGCAAACCTATTCATAT
AATGTTTTCTGTTATTGAATTTTCAAGAGACATGAGATCTGAAAAAATGTGTAAAGGCAAAAGGGAGGATATTTTGAA
TAATTACAAATGTCATTAAACATTTCCCTATTCTTGGAGGAAAACATTGTAAAGCAAATGATTAAGTGAAGCGGTGACT
TTAAGGAACTGAGACTACTTAATGATGCTAGGAGACTTCCTATTTGTATTTGTTAATGCAAAAAATTTTATCTTGGTG
GAAAGGCTCAAGCTTTCCAGATTAAGAGAACCTGAGTTGCCTACATTTGTCAAATGTAAACAGTAGAACTCTCATTTT
ACTGCTTCCTAGCTTCTTGAGAATCCACTGGCATAAAGAAAATGCTTCCACTTAATTAATGACTCTAAGCAGAAGTTA
ATGTTTATCAGAAAAAAGAGCCAGACTTATTGCCTAGTTAGAAGTTGTCACTTTAGGGCTATAAAATTTTATTTTGCT
CTGGTCTGAGCATATAACCCTCCAACGCTTACGTTTTTGGCATAATATAATCCAAAATTTGTATACTTAGAGGTAGATTT
TTCCTGGTTTTGATGAAGAAAGTTTAAAGAATTAGTTCTCTTTAGAATCTGAGGATGTTTATCTTTGGCATTCTGCAT
TTGGAAATCCAAAGAAAATGTTGGAATTTAAACAGTCTCTGTTCCCTCCCTACATGCCTCTCTTGTAGGTCTTGCATC
TTTCATTGTGAAAATTTAAGGCATACAAAAGAGGTAGAAAAAAGATCATCCTTAAACGATCACCTTACTTAGTGAGTTC
CTCTCATGAGAGGGGTTCCAGTGTGTCTTGGGAAGACCATTTAGTCACTCTTCAACTCAAACAATTCAGGCATAAGATGG
GTGGTTAAACTATGTGAGTGTCTGTTTCCCTACCAGTTATGAATTTCTATGATTCTATACCATGTTGTGCTCATTCGTA
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TACAATATACCATAGACTAGGTGGCTTAAACAGAAAAGTATTTTATCACAATTCCTGCAGGCTGGGAAGTTCAAGATCAA
AATGCCAACCAATTTTGTTCCTGGTAAGCTCTCTCGTCTGGTTTGCTGATGGCTTGCCGAAAGCTACTTTTTCCCTGG
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ACCTTCATGAATTCATCTAACACTAATTACCTCCCAAAGACCCACCTCCAAAACCATCACATTGGGGGTTAGGGCTT
CAACAAATGTATTTTCAAGGGGACACAAACATTTGGTCCATAAGAGAGTTTGATATTCTTTTTTTTTTTTTTTTCTGAGA
CGGAGTCTTGCTCTGTTGCCTAGGCTGGAGCGCAGTGGCGCGATCTCGGCTCACTGCAAGCTCCGCCCTCCCTGGTTCAC
GCCATTCTCCTGCCTCAGCCTCCCGAGTAGCAGGGACTACACGTGCCCGCCACCACGCTCGGCTAATTTTTTTTGGCATT
TTTAGTAGAGACAGGGTTTACCATGTTAGCCAGGATGGTCTCAATCTCCCGACCTCGTGATCCGCCCGCCTCGGCTTC
CCAAAGTGCTGGGATTACAGGCATGAGCCACGGCGCTGGAGTGCAATGCCGGGATCTCAGCTCACTGCAACCTCTGCCT
CCCAAGTTCAAGCCATTCTCCTGTCTCAGCCTCCTGAGTAGCTGGGATTAGAGGCATGCGCCATCACACCTGGCTAATT
TTGTCTTATTAGTAGAGACAGGGTTTACCATGTTGGTCAGGCTGGTCTCAGGTGAATTTGATATTCTTAAGGGATGAT
TGATTTAATAAGTCACTGTCTTGTTTAAGCCCAAAGGGTAGTGACTAGTATAATGGAATCTGTATGTTTTCCCAATTTG
GTAACACTGAAAATGATCTGGTCAACATCTTTCTCTTCATTTCTTATTTTCTAAATTTTATGTTAGGGACATTCTTACC

Fig. 9.93

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AATGATTTTTGAAGTCTAATAATCACTTCATAGACCTAGAAGCTAATTTTAATTTTACTGAAAATGATTTTCCCCTTTC
TTCACTATTCTTAGTTTGCTTATTTTATATTGTTCTTTCTTAACATCTAATCCAATGACAGGTCTCTGAAGTATCTTGT
CCTATCAATGGATTTACTATTTAACTTTTCAGATGTTTATATATTTTCAGAACTGACTATCTCAGTTACCCTTCTTCCCT
CTGCTAATGCCTGCTTTCTTACTGGCTTTTAGCTAATCAGGGGAGGGAAGGAAAGGTACAGACAAGAGAACAGGATATA
AAAATTTTTGTTGTAATTATATATTTTAAATCAGGAGTTAAAGTAAAAATTTTAGCATTTTCTTTTCTTTATAATATC
CCTGTTATTCTCCAACTCATGAAGTTTTATAAATCTTCTCATCTGGAGAATAAACTAGATAAATTTTAATTCTTATTC
ATTAATAAATTTTCTTGTGTATTAAATTCTAACTTTCTTGTAGTCTTTTGCATTAAATTCCCAATTTTCTAGTACACT
GAATCATTTTCTTCTACCAGTTTGAAGATATTCAAGATGTCATCTACTTAATTGGATTTGTTAATTTTCTTGATGAAA
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GGGTATCTGGCACATTCTTTCATAATGAATTTAAAGCCCACCTAAATAGGCTGTCTAGTTTTGGAGTCTGGGGGCTAGGAG
CCCATCTGGTGAAATTTTTCCCAGTAGGCTCCCTAGAATATTGCTTATTTTCATGGAAGACTTTAAAAAATTGTACAAA
CTTGGTAAACTAAAGTGATTTTATTCTTAAGAAAAATATATATTTTTTCCATCTGTACCTTTACATTGCTGAGTTTTTA
GAAGCAGTCACTCACCTAATGACCCAGGGTGAGTTATTTATGCCATTTTATCTGCATTTTTTTAGTGTCTTCTTGGAGTG
GGCAGCCAAAAATAAATTTAGAAAAGATGGTACATTTTGTATGGCACTTCAGAGCTATCTGCATTCATTTTTCTCTCA
CAATTACCCTGTAAGATAGAGTATTTTTTATTTGTATGGTTAAAGTAAAAATAACAAATTCCTTCCTAGATGGGCTTCA
ATTCTCTACTATATTCAACCCACCTAAGCTACGAGATTGACAAAGCTGTTGCAGCTTAGAATCCTAATAAAGTTAAGG
ATGTATAAACTGATTTTTTTTCACTGGTATTCACTCAGCAGGGTCCAAGAAGGTCACGGGAGATGCTGTAGGTAAATCC
TTTGAAAAGCACAAAATACAAAACAAAACCAAGTTCTGCTTGTGTCTTTATTCACAACAAATTGATATGAGAAGTTGT
AGTTTTTAGAGATTTCACTCCTTTGGTCTTTATACCTGCCTTCCAGCTTGTCTCTCTTCTTACAAAGTTCTACAAATAT
TACTGAATGGCTACTGTGTGTTAGGCTCTGAGTATCCAATGTAAGATACATGTTCTCTCTACCCTCATAGAGCATATC
TGATAACTAAATTTATAATTTTTTACAATTTGTTAAATTTATTTCAACCACCTGTCTCCCTAACCCTGATCACCTAACTG
TATCCCATTTTAATCTTTAATTCTTTTATAGGAAGGCTGAAGTTCTGTATCTCTCTGAAGTCTCTCTGTATTTGCACCTC
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AGTATAAATAAATGTGAGAATGACAGGAAAGGTCTGAAAGAACAGCGACTGATCAGCAACCATTTGCTTTAAATCAATC
ACATGTCACATGACTGTTTTAGTGGCCAAAGCCAATCATTAATTTGTCATCTTGGAAGGTTCCACTTTTTTCTGTAC
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TAAGTCTTAAAGCTTTCCTAAATGGAATTAGTCCATCAATAAGCAGATAACAACCTTTACTTTTATTGTTGAAAAATGTC
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TATAGTTAGAGAGACCCTGACCCTGCTTTCTTGTCTTCTCTATTACCAGGATATGTCATACGCCACACACATGGGGGT
AGGGATACCGCCCTACTTATAGCATAAGAACTGAGAATAAGACATTTAGCTTATAACTCTTTTATGGTAATATTCCCCT
CCCCTGCCATTCTTCTTACTCAGGTAGTCTGGACTTCTGTTTTGGCAATATTGCTTCCGTGAGAAGGATTTGACTGTA
CTTGCAGCCCTCAACATCCTGAATTTAATACAATCTACGATATTTGTTAAGCCCTCACCAACTATTTGACCAAACCTATG
ACAGTTACATCCCACGGAAGTGTATACCAAATGGTACCCACAGCAGTTTCAAGAGGTTAACACACCACCAATTCACCTCCA
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AACAAATGATAACACCACAGTACCACAACCTAAACAATCCTTTCTATAAATTAATTTAAACAAGAAACAACAAGCTTT
AAATGTCATCATCTCTGCTTATTGTATGTCCTTAAAAATATTATTAAATGTCCAACCTTTTATTTTTCTAGAAGAGGTCA
TTATATAGCATTGATTTGCCAGCAGGGTTCTATTTGAACATACCAAGAGACCTAGACATTGCTCAGAAACAGTAGTCTC
AAAATAACAAGGGATTGGAGGAAAAGATGAGAGATCTCCAGTATGTCTGCATATAAGGGCTGAAAAAGTAAAAGTTTC
CAATTGTTTTTTTTTCTTTGCAAGTGCTCATGCAGGGATTGAGAACATGCATTCCTCACCCAAACATAAATGAAATAATT
GGCAAGTAGTCAAAGAAGACCATATCCTTGAGTGGGTAGTATTGTTGTCTATTTGGAAAGCATTTACATGTTTTGATTT
CCTGAAACAATGCTGAAAATGTCCTAATGCAGGAAGGGAGAATTGAAAACAACCACCATAAAATGCAATTAGATGTTGG
TAAAAGTGAAGTCAAGATAGACGTAAGTTAAAACCTTTCTGACAGGGTTTTCTAGCACTGGGACAGTATTTTTTTGAGGAAA
TTATAAAATGGTCTTATTTCAAGATCTTTTAAACAATCTGATGAAATATTTATGGTTCTTTAAATTTGTATTGTTAGTAT
CATACTCTGAATTATAAAACATTTAAACATTTTAGTTTATAATCTTTAACTTCTCACTATTATTTATAAATGTGTAT
ATAATTTGTGCATATGTAGACATTCATGAGGAAGATGAACATATATGTTAATTGGCATCTGCTCATTTTAAACATAAAGT
TGTATACTTTTCAATTACAGTAATACACGTCATTCATTAAATTATCTTGTGGCTTAGCTTTACAAATTTTACCCTTACAT
GACTTTGGGTGAATGACCTCACCGGTGTCTATATCACTCATGTGTAACATGAGAGGAGTAGATTAAATAAACGAAAAGA
TTTTCTTTCAACTCCAAAGCTATGACAATGTATTTTCAAGATTGTGTACTTCTTAGAACAGGCTCAATAATAATTTTTCA
CATTATGAACCTTTGCAGTCAAAGAATAGGTTCTTCTTAACCTAACAAATGACTATCCTTTCCACCCAAAGTATAAACAG
GATCCTGATTAATAAATTGAGTTCAAAGAATCTCTGCCAAGTATGCAAAATCACTGCCTCTGTTTGTGCGACATTCATT
GCCTCTAAGAATTGATGGAATTGAAAATAACCTCATTTTACTGGGACCTCAGAGAATTAATTATTTAAATTTTTTGCTG

Fig. 9.94

CTTTAAACATTAATTTTCTTAATTTACCCATATATGTTGCTGATAAGAGCTGTAATATTTTGAATGGTTGTGCTTTGAA
GAAATCTGAATCCTTTTGTCTTTGTATTCCAATGACAGCAGCTTTGACCAGCGACCAGCTCTCTTCTGAAAACCTACCATT
TTGACCCCTGCATTTACCTGTTGTCCCACCAAGCCTGTCTGTTTTTCCATTAACAGAGACCTCTGAATTTGTTGTGCCA
AATGTGGACAGTGTGTTTCTTCTCACTTTCTCTGAATTATAACCAGTTCCAGGCGGTAACATGCAACCGAACTTTACT
GAGTCTTGACATAATGGGAAAAAATCATAGTGTAGAAAGAAAACCTATAGGACTGAATATAAAAAAATAATCATTTCTG
GCATTACAGGACAAACCCAGTCCTTTCTGCTTAGTTACTGACCTACCCCTGTTGCTTTGCTTATCTTCCCACAGTGAA
ATGTCTTTCTTCTTATATCCTACATGGTTTCCAGGCCCTTTACTCCAGGAAAGCCAGGAGAAACGCCTTATTCCAAGTT
CAAGTAAACATAATAATTTACAAAGATACAACCTCTGCCCACAACAAAAACTCCTTTTACAGCGTTATGCAAAGGCATT
TAGACTGGAACATCTATGTTCCAGACACAGACCTTAACCAGTCTTTTGTCAAACATAAAGAGCAATCTTCTCTCAAAGC
TGGAATAACACCTTTTCTTTTAAATAACATTTCTGTGTCTCACACTCCCAGATGTTTTTCATTTAAGACTTTAGAAAATA
CTGGGATCAGTTATCAGCCAAGAGTACCCCATTTCTAATAAAAAATATTTAAAGACATGGAAAAATCAATGAATCCAAAC
AATCATCATCCTCACCAAACCTTATCATTTCTATAACTCACAGTAAATAATCTCAAGTTCTTTATTTTGGTAAATTAA
GAAATTCAGAGTAAACTCTCTAGCTTCTGATTTAAGCTCAGAGATGCAGAGAGCTTCAGAGTGTCTGTTCTCATTTCTTA
CCATAAGGAAAAATCTATACAAACTACCTATTCTAATTTAAAAAAAAGCACCAGCAAATTTAGGTCGTAGGGCAACC
AAACAACTCCAAATCTGGAAAGAGACAGGCACCTGCAAGAAGAAAGAGGATGGCATCATTTGTTTTTCTTGGGTAGAC
ACCACCAGATGTCATGTAAACCAGCAAGATGATTCAGCTAAACATTTTAAATGAATTGCTAAGGCTGAGTATGGGCTAGC
ATGAAAATGTGACACACTGGAGGTGGCAGATATAGGGAGTGCATCCTATAGCAGGCTTTTCTCCACAAACCCACCAG
GCACTCACAGGAAAGACTGGGGAGAACAGCAGCCACCTTCAAACCCACAACCATTTCCCAGTGGAACAAAAGAGTTAAT
TGGCAAAGAGAATAGCAAAAATCATTTGTCTTAGGGAACTGGAGGAAACCCATTGGTGATGGTGGCAGTAGGAAGAATGA
TCGTGGTGAGGGGAAGAGAAAAAAGTAATGCTCTATCCCCAGGGGTGGGGTATGGAATATATGCTAGGATTTGCACAAC
AATTGGAGAAAGTTGCAGGAGCACTTGTGAAGGCCACATTGCTGATACAGAGGTACACATTACCTACCTAAGACTGAGT
CTTAATCAGAACATCAAGGAATGCCCTTCTCCCTGGCTGTACCAACAGCCTAACAAGTGTGAGTAAAAATAATAT
GGGAATATGGTTGAATATGGAAGAAATTCAGAGACACATTCTCTTTAGGGCCCAGCATTAAGGGAAGACCCAAAGCTA
AAGGGGGAGCAAATATTAAGAAAATAACAACCTGGCAAGCCATTTACAATCTATTCTCTTTAAGAATCCAAAAGTATC
TATCTCAGTATCTACTGTCTACACAAGATATCCGGCTTTCAGCAAAATATTATGACCATATGAAAAGGCAAGAGAAAG
CACTCCGAAGAGATAATACACATAAACATGTGATATATGGGACACATATAAAAATTATCACACAAGGAATTTAAAGTAA
CTATGATTAATATTTTAAAGGTTCTAAAGGAAAAGGTTGACCACATGTGAGATCAGATAGGTAATTTTCAGAGGAGATAT
GGAAACTAAGAAAAATCAAATGGAAATGCTAGAAATTA AAAACATAGTCAAAGAGATTAAAAAATGTCTTTGGGCTTGT
TAGTAGACGTGAGACAGCTAAGGAAATAATCACTGGACTTGAAAATAAGTCAGTAGAAATTACCCAAACAGAAAAAGA
GTGGAAATAAAAGGGAGAAAGAAAGAACATAACAGAGCATCCAAGAACTCTGGTACAATATTAAGTGGCATGACATATC
CATAATTGAAATGACAGAAGGAGAGGAAAGAGAAAAAGAGGCAGAAGAAATGTTTGAAGAAAAAAATCGCTGAAAAAAT
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ACACACGTGCAGAGTCATGTTATATTGAAATTGCTAAAAATCAACAACATAGAGAAAACCTTAAAGGCAACCAGAGGGA
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AAAAAGGAATGACATCTTTAAAGTATTGAAAGAAAAACAAAACCCAGTAACCCAGAATTCTATGCCCAATAATAATACA
GTTAAAACAGGAATTTAAATTTTTCTGACTGAACTTTTTAGGTTCCCTATTCTGTCTGATGTCCCCAAAGTAATATGCAC
CTATATCCTCATTTTATTTTGTCTTTCTTTTCAAATCTGAGAGGGTAAAGCTGTTTCTTCTATTCAAGACCAACACC
TCTATTGCCTCAGAGGGTTAGTCAATTGCCTCCTCCCTCTATAGGCTCAATTGCAATTTAGCTAATCTCTCTTCTATTT
CTCCTTTTTACCTTAGTTTATGAACCTAGACCATCCAGTCCTTTCTGTCACTCCCGTCCATGTATATCTCTTTTGACAAT
TCTAGTATTTATATTTTTCTTTCTAGTAGTTATTGCTATTTCTTCTGAGACTGCATGTTCTTAAACAATGCTGTTCC
CCAAGATTCAGTCATTGGCCCATGTAATATTTTAAAGGTATGAAGACACCTGAAAGATCATAAGATATCTCCTTTCATGT
CTTAACCTATCTTCTATTTGATTATCTCTCAATTTCCATATTTAGTTCCCATTTCTACTTTTAGTCACAGAACTCTAACC
AAGTGCTTGCTGGACATTCAACATCTACACAACATTTCCCAAACCTAGATGCATCCATTATTCTTCTCATCTTTAAA
CATACTGCTACATATCCTAAATATTTTCTATTATATTTACTGCCTCAGCTATCCAGTTGCCTAAGGCAGAGCCCTGGGA
ACCTTCTCAGCTCCAGACTCATCATCCCTCCCAACATCTAAACAAGCAACAAGTGTACTGCATGTATTTGTGGGCAC
ATATTTCTGAGTCCTCATCTCAGCTTGACTTCCCTTTCCACCAATGCCCTGGGGATCACCTTATTTTTTCTCCCCAAGA
TGGTTCTCATTAATAATATACCACAGTAGTGATAAGTAGTAAATTATATATGGAAGCTGCTGACTGTTGCTCCATTAGC
TGTTTCTGTGGCAACTCTATTTTTTCTTATTAATAGTGTACTAAGTGTGTTTGAATCTATGTCTTTTCACAACTTCTTTT
GATTACAGTTTAAAGATAGCCTTTAAATACTAAATTCATATTAAGGTGATCTGATTGTGTGTGTGTGTGTGTGTGTAATTT
CTGTTACAGCCATCATTCATAATTCATTATTTGTGAGGAATAATTATAGCAAAGACTGATTAGATGAAATGTTAGATTA
TTTATTATACCAGATTGGAATGGAATTAAGTCTAATATATGTGTTAATGTGACTAATAAGAATAACAATAATAATTTA
GAATTGCAAAATGCTTTTAAAGTTTCCATATACTATTACAACATTTTCTTGTTTTTTTTCTTTTATAAAAACTTACAGGTA
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GGTGAATCCTAGATATACTTCAGTTCTTCTGATTCCAAATTCCTTACTTTGGACACTAAACCTTGGAGCCACACAAATG
TAACTGTAGGCATCACTTATGTACTAATGTACCTGTTACACCATGTGCCTACTGGAGAGAACTGTAAACATATTTATAG
TCAATTATCTATATTTATAGCTATCCTCACAGTTGGATAACAATGTTATATTGTAAACTCTAGATACTTTTGCACCTCTA
TGACAGAAAGTGGATTTTTTGTTTAGGAAGACATTTTAAATTATGAATGGAGGCTTGTTGGGGATCCCTATTAGCCTTTG
TTAAATCACAAGGAAGGTAACTCTTTTGCATAATAAAGTATTAGGATTTTTTAAACTTGTCCCCTGACTGCTCAAGAGA
TGAAGTTTGGGGAGCGGCAGTACTTTGGAGGACTCTGGTAAGTGAATAGATATAAGAAAGTTCTGCAGGGATAGAGCCA
CATGGCAGCAGACAGGGCTTAGGAAGAGGGACTACCAATTGTCCCTGTAGCAAAGATGGCCTGGTTAAAGCCTCTTTAG
AATGAATGAGTTCTAGTAACCAGTTTTTAAAGATTTGTCTGTTTCTTTTCTCCCTCAACCTTTCTTTAAACTTTCTGT

Fig. 9.95

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AATCACAAACAGCATTGTAACAATTTTGCCTCAAGCTGGGAAGAGAAGGGCTGTGTTCTCCTAGAGACAAAACAGGGAA
GAAAGGTAGCAGGGGCAGAGTGAAAGCAATGATTGCTGTGGAATTGGAACAACTGTGCAGAGGAAGCTGTGACAAATAA
TTAGGGTTGGTGAAGCACCTGTCCCTTTGGAGGTATTTCCCAGAAATACTGGGAAGGGCTCTAAATGTCCCATTTGTTAT
GTGGGGAGATGGTTTTTGTATTATTGTATTGCCACAGAAGTCTGAATACGTTCAAATATTTTGCATATTTGTGTGTGTGT
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TATTATATATATAATATAATATATATATATTTCACTCATTTAGTTCTCAAACTATTCTATGACACTAATATTTTATGTA
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AAAATACCTAAAACATACATATTTGAAATAAGAGAATATGAGGAAAAAGTCTCAAAATTTTATGTAGGTTTAAATAATA
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GATTAGAGGATGGTCATTGAGATTTCACTGGATTTAAATCCACAGTGGGATAGGTTTAAATCCTTTCTGGAAAAAATAT
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CTCAGATCCATGCATTTCTCTAAGTTTATACTTTGTTTAAATTGAGCTTTCATTGTTTCTATGACCAATTTTCAATTT
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TCTCTCTGTCTCCAAAACAATTGAAATTCTCTCATATGGTTTATTGCCTTGCAATTTACAAAGGAGCCACAAAGTTCTGA
TTTGTGTATACTATTTTGTCTTAACTAGCTATCTGGCTGATGTGCACATCAACAAATGACAATGTAGTCATTCCATCTT
TGGTACATGGAGTATTATTTGATAAAAATTCCTATATTTTAACTTCTGAAAGTAAGGTGATTTTGAAGTATCTAGAA
GATAGTTTCTTTATTTCAACAATCATAACCCTGTGCTGCCAGATACATATTTTGATCCCAAACCTTGAAAATATTTCAAT
GGTTAGATTATTATGCTTTTCTATCTGACAGATTTTATGGTTTACCATTTCCTTAAAGCTTTCCAGCTTTTCTCTCTC
TTTAAAGTAAGTATTTGGAAGTTTCATCATTTCCATTATCAATACTAGAAATTAAGAGTCAGAGATATATGTATTCT
CAGAATTGTCTGAAGAGTTTATTGTAATTTAATAAGATGTTCTTCTGTTGTTTCATCTATTATGTATTACATATCA
TCTATGTCATTATCGTGTCTCTCTCATGATTTTCTGGATCACTTTAATGCTCTAATCAAGTGCTCTTTATTTTGTGT
ACAGAGGCCTCATTATGATAATACTATATAAACATTTGTATGATGCTTTTCTAGTTTACAAAGTCCCTTAAAAAACAT
TACCTTATTTGCAACTCATAACAACCTTGTACATTAGGCTGTGTTAFTATTATTATTATCTGTAGACAAGAAAACACTT
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CGTTATTTCTTCATCTCTTGCCTTTATTGAGCTTTTCTCTTTGTCCATTCTACAGTAGGAATGAATTCTATCTCATAGA
TAGATTTAGAAACCAACTCCAGCAAAAAAATTTGTTGTGCTCTTTGTGAGGGAAGAGGTTAGATTAGGATGAATTAAATA
TTTCAGTTGAAAAAATCTGAGCAAATTCATTTAAGATATTTTGAAAACCTCCTAAACAGTATATAAATATATGAAAGTT
TAAACAATAATAACAGTGGAGGTACATCTTTTGTGTTGTAAAAAATGTAGTTGATCTTTTTCTTTAATGTTTCTCTTAT
TTTTGTGTTTTGCTTAGTGAGGCTTTTCTACTGCAAGATTATGGATAATTTTGCCCATGTTTTTGAACATTCTTATGGT
TTTTCTCTTTTCCCCTAACCTTGACAAACAACTTCCTTTCAAATGTCCTAGATGCTTCAGGGATAGCTATTTGAAATA
GTTCTTGGAGATTAATAATTCTCCATGGCAAGCTGTCTTATCCGTATTCTGGTAGGTTATCTTATAGAAGAGGAAAAAGA
AGGCAAAATCCTAGAGTTAAAAAATACAATAAACAAATGAAAAAGTAGTAATAACAAGTACAACAACAAGAGCTCTA
GTGGTATACTGAGAAACAACTGCATTTCTTTGTTCTTTTTTCTTATTTTCAATCATTTATGTTGCATGAGGTACAGCCAT
ATGAATAAATATGTACTAACACATACCTAGACATACACAATCAGCCCTCATATTCTGATTTCTGCTATCTGTGGTTTCA
GTCAATTGCAAATCTAAAATACTTTTTTAATCATCTGTAGTAAGCATGTAGAGACTTTCCTTGTCTATTATTTCTTAAAA
AAATACAGTGTAAGTATTTACATAGTATTTACATTGTATTAGATAATGTAAGTAATCTTGAGGTGGTTTAAAGTACACA
AGAGGATGTGCGTAGGTTACATGCAAATACTATGCCATTTTGTATCAGGGACATGAGCATCCGTGGTATTCACTGGAAG
TTGTGAAACCAATCTCCACGGATACCAAGGGACAACCTATATACATATATGTGGCTATGCGGCCTACCTACTCAGAAGA
GGGTTACAGAGAAGAGGGGCAGCATGGAGAGATTCTAATGTCTTGTGAGGGAGAGGGGAGCCTACAGTGGAAATGGTTA
GTTAGAAAAATAGCCAATAACTTATTTCCAGTGAGGAATTAGAAAACAGGGATTATGGAATTGAAAAACTGAGGCATTA
AAAGTCTTCTCTCTGAAATTACTATGCCAAGTGGAAAACTGTCTCTGAACCCCTATCCAGGAAAACATTCTTCTTCAAA
AGATGTTACTTTTAACTTTGATAAGTCTTGATTAGCTCTTTTGGGTTTTAGCTGTTCTTTCACTATGTCCACACTTTAC
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TACCTGATTGCACAGGTAGCAATCCAGGCTAAGAGTACCCCTTAATGCTGCAATGGCCACAGTAGTCTTGGACAGGGG
GATCATGCCAGTAGGTCTGCCCAGAATCTCTGGATAGATTTATTGTTGAAGACCATTCCTAGACAAATCCATTCTGTAA
AGACTGGAATAAGTATTTACTTCTTTAGATGTGCAGATATTGATACATGTCCTCAAACATCAGATATAATCAGGGAAAC
ATGATGACACCAATGGACAGAATAAAGCACCAGTGATTGACCTTAAAGAGATGGAGATGCATGAACTTCTTGACAGAG

Fig. 9.96

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AATTCAAATAACTGTTTGAGAAAGCTCAGTGAATGTTGAGAAAGACAGATAACAATTTCTTTTAAAAAAGGAGAAC
ATTAAGTGACCCACAATGAGAAACAGAAAAATTGAAATAACAATTTTAAAAATTAAACAGAAATCCTAGAGCTAAAATA
TACAATGAACAAAATGAAAATGAAATAGCATCAACAGCAGACTTGATCAAGCAGAAAAAGAATCTGTAAACTTAAACAC
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CGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCGGGCGAGGTGGCGGGCGCCTGTAGTCCCAGCTACTCGG
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CTGGGCGACAGCGAGACTCCGTCTCAAAAAAAAAAAAAAAAAAAAAAGAAAAATACACAGTTCAGGAGAGGAAAGAATGA
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CAGGTCAGGAAGGTCAAAGATTTCCAATCAGATTCAGTTAAAATAAGACTATTCAATACATATTACGATAAAATTCTCA
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TATCAGCAAACCTTCTCAGCAGAAACCTTACAGATCAGGAGGGACTTGGTGATATATTAAAAAGTGCTGAAGGAAAAAAA
AATGCCAGCCAAGAATGCTATACACAGAAAAGCCATCTTTTAGAAATAAAATACTTTCCAGAAAAATCAAAGCTGAAGA
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CAAAAACATGTGTTAATGCTAGTAACACAAAACATCTATAAATATAAAACTCATTGGTAAAAGTACATAGTCAAATTTA
GAATACTCTAATATTGTAATGGTGGTGTGTTAAATCACTTATATCTTTAGAAAGAAAGGTTAAAAGACTAAATTAGTAAAA
ATAATAACTACAATAATTTGTTACAGGACATGCAGTATAATAAGATGTAAATTTGTGACACCAAATTCAAAATGTGTTT
GGGAGAATGAGGTAAAAGTTTAGAGTTTTTTTAAATTTTTTATTTTGCAATCCATGTTAAGTTGTTATCAGCTTAAAAAAC
CTGTTAAAAGTAAAAGTGCTTTTTATAAGCCTCATGATAACTACAATGGAAAAATAACTTGTTAAAATTATGGAAACCT
CGTCTCTACTAAGAATACAAAAAATTAGACAGGCGTGGTGGCGGGCGCCTGTAGTCCCAGCTACTCAGAAGGCTGAGGC
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GACCCTATTATTGGTACACACAACACAAAATGCAAGGAATCAGAATACACTACTAGAGAAAATCACTTAACCACAAAGA
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AATCATATTCTGCCCTCAAGGGACCCACTTCACCTGTAAGTACACACATAGATCAAAGGTTAAATGTTATTATATATA
TATAATTATCTATAATATAAAAAACATATTTATATATAAATAATGTTATACAGATATAAAATTGAACACACAAATAAATAG
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AAGCTAAATTAGCCTAAGCAGCGTGATATCAGCATAAAAAACAAACAAACAGATAGACCAATGGAACAGAATACAGAGCC
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CATAAAGAAGTCTCAATATGGTTAACTAGTTTCCAGAGCAGTGCTACAACCTGAGCCTTGTGCATCCCTGAAGTGATGAG
CACAAGTATGATAATCATCGGAAGAGAACATATATTGTATAATTTGGAATCAGCCAGTTATTTGTCAGACTACCTTTGC
TGTCTGTAGGATCAGACACACATGTGCCTCTGCATTTGGGTAAATGTAAAACCAATTTCTTATTATAAAGAGGAAAGA
CTGCTGGAAACAGCTGCTCTGGAAACCAAGTTGCTCAGAGGAAGTGAGCTAACTTGTGTTTAGCTAAATGGTGTGTAG
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AGGCAAGAAAAATAGGGATAGAAATGGAAAAAGTGGAGAGAGAATAACAATGGGAAGGGGAAAGAGGAAACACGCTTATG
TGTTGTAGTGACTGACCAACGCAGTTTATGAAACAACATAGCTAGCTCAGGAGTTAGCCCAGGCTTTCAATTAAAAAGA
GATGCATAAGCCCATATACCTCTTTTTCTTTCCCTCAAATGACCTTGAGTTTTAGTTTTGGTAGGGTTTTGGTACAGCT
GTTATTTGATAATATCACATTGTGCTTTTCAATTAACCTTAGATTGTTTTAAAAAACAGATAACTGAAAACCATCTGTTTG

Fig. 9.97

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TGGTTCTGTGTAATTTGTTCTCAGGACAGGGACTAGGAATGAACCATTTTAAATTCTGCTAATGAAACCTCTCATTAAT
TTAAGGAGTTCTACAACTAAGTCTGTGGTACCAGGTAGAAGGGGTGCCAAGTGTGGGGCTTTCTGGGATAAGGGGAAAA
TTAATTTCTTCCTTTGCCATTTGTGGTTGGGTAGCATCTTTTTTTTTTCTTCCTGGCAGCGAGTCTCATAATTTAATAT
GCATTAACATAATCAGGAATGATTGCTTAAAATACATGTTCTTGAAGTTTATCCCAAGAGATTTTGTATTGAGTATCTC
TGGGAAGGGCTCCAGGAATTGCCACAGTTAAACAAGCTCCCCAGGTCATTCTGATGTAGATGGTCTGAAGGCCACACTG
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TGTTTTCCCTCGAGCTTATTAAGGGCATAATATCACTTTCCCTCTTGTTAAGAGTAAAGATTGGAAGAATTTAGTTATC
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TCATAAAGTTGTAGCATAACATATTACTCATATGTTTATGGTGATGCAGGTGCAACAACTCACTGCATGGCCGGTGA
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CGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGGCCGAGGCGGGCGGATCACGAGGTGAGGAGATCCAGACCATTCT
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CCACTGCACTCCAGCCTGGGCGACAGGGCGAGACTTTGTCTCAAAAAAATAAATAAAAAATAAATAAATTTAGCATAGC
TTAAGAGTACAGTGTTTATAAAGTCTACATTAGTAGACAGCAATGTCCCAGGCATACTCACCCTCACTCATTGACTCA
CCCAGAGCAACTTCCAATCTTGTAGCCTCCATTCACGGTAGGTGTTTTCTACAGGTATATTTTTTATCTTTTACCACATT
TTTACTGTATCTTTCTTATTTTTTATATGTTTAGATACACAAATACTTACCATTGTGTTATGCTTGCCTGCAGTATTCA
GTACAGTAACATGGCATAACAGGTTTATAGCCTAGGAGAAACAGGCTATACAATATAGTCTAGGTGTGTGTGTGTAGGC
TATACCATCTAGGTTTGTGTAGGTATACTCTGTGATGTTGACACAACAACGTAATCACTTATGATGGATATCTTAGAAC

Fig. 9.98

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GAGGTGATATTGTGGTTTTGATTTGCATTTCCCTGATGATTAGTGATGTTGAACATTTTTTATCTACATGTTAGCCATT
TGCATGTCTTCTTTTGAGAAATGTCTATTCAAATCCTTTGTCAATATTTTAATAGGGTTATGTGTTTTCTTGATACTGA
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GGGTTGCCTTTTCACTGTGTTGTTTCTTTCGCTTCCTGGTTGCATTTTGTGTTGATGTAGTCCCCTTGTCTAGTTTCA
CTTTTGTGTTTGTGCTTTTGTGATGTCATATCCAAGAAATTATTGTCAAGACTAAGAAAAAGAGAGAAAACCTCAAATAA
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CTGGGACTACAGGCGCCTGCCACCACGCCAGGCTAATTTTTTGTATTTTGTAGTAGAGATGGGGTTTACCCTATGTTAGGC
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GAAATTAAAGAAGATGCAAATAAATTGAAAGACGTTTTTGTGTTTATGGATGGGAAGACTTAATACTGTTAAATGTCCA
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AGTCCATGATCTCTTACTATTTTACTTTCAGCCCTGTAAGATTCATGGTTGTATTTCTTCCATATAGAACTATGGAATA
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TATATAATTCTATATACTAGTCATATATACTAGTATATAGAAATTATATATACTATATGTATGTATATATACACTAATAT
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CAAAGTCTTTCATTAAATGATACCAGAACATAGGATCTACTGAAATTCATAATGGCCAGTTGGAAGAGGAAGGTGATGT
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CACAGTTTCTTAAACATAGATATCTCTCATATATTATCAAACACTTATGGTGAAACACTACATTTTTTGTTCATAGGAGT
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TATTCAAAACCTCTTTGAATTTCTGGTTAAATAAAAAATGCCAACTATGCCTTTTTTCATTAAATAAGTCTATCTTTCTATTA
TCTAAATGTATGTCTACTATAAGTTCCCAATGGCAGCTATGTCTTGTGCTTAGGGATTGCAAGATGAAGGTTAACTA
TTAACGACAGTGTCTTCTGAACCTGAAGTTATTAGAAAATCTTTAGGGACTCCTTGGACTCTTGGAAATTATGTACAAA
TTATAATGACTCATTTGTACAGTCATCCCTTGGAACTGTAGGGGATTGATTCCAGAGCCAAATACCAAGACTGCAGAT

Fig. 9.100

ACAAAACCCAGTGGATGCTTAAGTTCCTTACATAAAATGGTATAGTGTTAATATTTAACCTATATTCTTCCATACACTT
TATCTCTAGAATACTGCTAATAACTAATAATATGTAAATTCTATGTAAATAGTTGTTATAACAATGATTTTTTATTATT
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TTTCTGGTATTCAGATTTTTGGGTCTATCTCCTTTCAAATTGTATCAGGGTTGATCTCTGTGGCCAAGAAAATTCAGGAG
GAGTTATGATGTGCCACTTTGGAGGTTAGGTTATAAAAAGTGTGGCTTCTGTCTTGGTTACAGTTTCTCTTTCTCCAAT
CACTCACTCTGGAGGCAGCTAGCTGTCAACTCACAAAGACACTCAAGCAGCCTATGGAAGAAGGCCACATGGTAAATA
TGGAGGCCTCCAGCCAACAGTCAGCAAGGAAGTGAACAACCATGTGAGTGACTCGAGAAGTGCTTCTCTA
GCTCCAGTTGAGACTTGCAGTAGCAGCAGCCTCAGCTGGCGGCTTGACTGCAATCTCTTGAGAGACCCTAAGCTCTCCT
GAATTCCTTGATCCTTAGAACTGTGTGAGGTAAGAGATATTTGTTGCTTTAAGATGCTACATTTGGGGATAATTCTTA
CACAGAAATAGATATCTCATTACATTATCTTGACTGGTTCATGATTAAAAGAAAAGTGAATGTAAGAAAATAAAGTGTT
TTAATGCTGACCTTCCCTGTTAATCCTAGAAAATTAGAGTTTGAAATAATAATGTCATAGTCACTATTCCTTTAATCT
TGTTGGTAAATAATGAATGCAGCGTGGCCCATTCACCGCCAGCACTTGGTCACCATTGTGATCTACACAGCAAGAAGCA
GCCTAACGACCTGTCTGTTGAAACAAACAAGTTTTCTTTTAAGTGATTTCTTTGTTTCCATTTATAAGGCACCAACTTT
CAAAGGTGTTCTGGGAAACCTTTCTTGATTTCTTCTAAGCAAAATCAATTCAACAGAGAGTTCACGCTTTGCCTGTGA
TCAATGGGGAAGTACCAGTGATAGCTTTCTTTTTTTTTCTATAGGGCTGCTCATAGTCTCCTCCGATAGACTTTACAGCTGT
TAGTTTTGCTGCAGTAGTGACTTGCTAAAATGGTGGCTCATTTGAATGGTGCTTGTATTAATTTACACTCCCACCAACA
GGGTACAAGGGTTCCCTCTTCTCCATATCCTCCCCAATATTTATTATCTTTTTTTCTTCTTCATAATAGCCATTCAACA
GGTATGAGGTGATATTGTGGTTTTGATTTGCATTTTCTATGATTAGTGATGTTGACCCTTTTTCATATATCTGTTGG
CCATTTGTATGTCTTTTAAGAAATGTCTATTCGATCCTTTACCTTTTTAAAAAATGAGTTATTTATTTCTTGCTATT
GAGTTCGTATTTTGAATATTAACCTCATATCAGGTGCGAATATTCATATCGGATGCAAATATTTTTTGTATTGTATA
GGCTGTTTCTTCATACTGTTGGTTATTTTCTTGTTGCTGAGATGTAAGCTTTTAGTTCGATGCATTCCCATTATGTAC
ATTTGCTTTTGTCTTAAATACACCTTTGTATTGAGTAGATCAGATCCTTTATGAAAGCAAGAAGGGATCAATGATTAC
CTGATGGAGAAGAAGAAAATTGTGAGGAAAAGGGTCAATGAGGACATTCTTTGCTGTTTTGCAATTTTCCAATGAGCTG
GTCTTGATATTATGTTTGCCTAAATAGCTTGGTCTGATCTTGGACACTAAATCCAATCCAGCAGGTTCTATCTGGAAG
GAGACTATATGATGGTGGAGACACAAGTCTGAAGGTGAAATATTGCACGATGTAGAAATAGATCAACCATGCACATTTA
TAGAACATCTACTACCAGGTGCTAGGTTCTGTGCTGAGCAATAAACATAATTCCTCTGTAATTTTTGTTTGTGTTT
GTTTTTGAGATGGAGTCTCACTTTGTTGCCAGGCTGGAGTGCAGTGGCGCGAGCTTGGCTCACTGCAACCTCCGCCTC
TCAGGCTCAAGCGATTCTCATGTCTCAGCCTCCTGAGCAGCTGGGACTATAGGCGCACGCCACCACACCTGGCTAATTT
TTTGTATTTTAGTGAGACGGGGTTTACCGTGTTGCACAGGGTGGGCTTGAACACTGAGCTCAGGTGACCCGCCTGC
CTCAGCTTCCCAAAGTGCTGAGATTACAGGCGTGCGCCACCACACCAGACTCCTCAGTAATTTATAACCTAGTTGAAA
GATTTGGACATATGTATCACAAGATACCTATGAACAAAGGCAAGGGCATATCTTATTTATAAGATGTTTAGATAGATTT
CAAAGGAAGTTTTTCAGCCAAAATTCATCATGTTTCCATTTTTGAACTAAATACCAACATATCTCAAATTCAGATC
TCAATTACTTCTACTGGCAAAATGTAAAGACCATTCCCTCATTTGTTATTTCTCCAAGTATCCTGAGTACTGGAT
ATCTGAAAGGCTCTCTGCTAGGTACTTTGTAGGGTAGGAAAACCTCAAAGGATGTGTGGAATGAAAAAGTTTCCAAA
TACTTCTTCAAATTTTGTGAACTTATCAACTGAAACAAGGTTGATTTGGGTCAAACAAGGTTTGTGAAGTAAAGCAA
AGCAATTTTAAATAGGCTAAAACCCCATGGGTTGCAATGAAAGCAATGTTAAGATGACTCTTAAGTAACCTAGAACTGT
TGAGGACTTTGGTAATTAATACTCTTCTCATAGCTCCCCCAGCAATCAGGAAACAAGGATAGTTTGGATTAAGGTCT
CATAAATAAACATACAGCTTGTGAGTCTTGGAGATGGAGAAGTCATTCTTGAGTGTGCCACATGGAGGGCTCAGAGAT
AAGTGATGACCCTCTTGTCACTCTGGCAGAAAATATATTAACCTCTTTAAACCAAGAATTATAGTGATTTCAAGTGGTAC
AATTACAAGGAGTGACTTCTGGGACTTGCATAAATCAAGCTTATTTAAATAGTGTTTCAGAAAAGAACATATGCTACTA
TTAAAGAGTTGGAAAAAACAACCTGCAATCCCATAAGATAAAGTTAGGTGAGAATATTCATATTTGTTTCAGATACAAG
TCCATTATTTATGTCAAATAATTAATACTCTTTAGATATCCATTCCAGGGGAGGCTCTTGAGCCTTTACCTCTCTC
ATGTCCAAACTTACTAGTTATTTCAATACTGAAATAATTCAGTTGTTTCATGAATTATTTGATGAGATTTAGAGTGATC
TATATTAGTTAATTCATTTATCAGATTTGTTGAATACCTGTTATGTGCACTGGATGGCAATTGGTGCCATGTTGTATAT
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CCAAAGCTGAGAGAGGAGAGAAAGATTATTTAAGTTGGTATGGTCAGATAAGTAAAACAGTATTTTAACTAAGCTTTGA
GAAATAGATAGAATTCTACTTGGAACATTGAGGAGGACTCCATGGAAAAGTAATAAATGAATGAAGACCAGAGCCAGG
CTGGAAAAAATCTTTTTAATGGAATAGCAGTGAGCAATGCAGAAAAGCTAGAGATCATCTTACTGCTGTGCACTCTAA
ATATTCCCCATGGCTGCTACCTTGTCCAATTCCCTCACAGTCTGCTGCTACAGAAAGCGTAGCAGTTTTCTCTTTAT
CGATGATGAGCAATAGAAAGAAATGCTTCACTCTGCTTAGGCAACAGGCACATTTAATATGGGATTTCTGTTGACAGTG
CCTTAAGCCATGGAGCTAGCTAATCAGCACCTTACTCATTTGATGACAGGAAAGAGAAGAGATCTGGGAATGAGCAGAT
CATTCAGCATTCCTGCAACATGGAGAAACAGAGCCAGACAGCTAGCTCTCCCCTCCAGTTTCTATGAGTTCCAATGT
GACCAAAGCCTTATCTTGGCCTTTGCACCTGGAGGTACCGCTGCCGCTGGGAGGATGAGTGGAGGAACTTGGGATGT
ACATGATAGGGTCAGCCAGCCAGCAAATGAGTGTTTGCAGAGTGACTAGGGAGGAGCAAAGGAGAGCTTTCAGTGTGT
AATAGTCCAGAAGTTGCTGCTGAATCAAGCGAAAAGAATAAGCAAAGTGTTTCAGAAGTCAGCCTGCTTCACATCTTGT
TGTTTCAATTTATTCCTCTATTCTGACCACTGAGCCCACCAAGCTCTCCTTCTTGATATAACTTTTTATCATATTTAATAT
TAGGGCAAGACTAGTTTGGGATCTTATGTGTCTAATTTATATAAATGTAGCAAATAGTCTTTTTTGATGGTAAGATAAAC
TAGTGAATTATCTTAACATAATCAATAAATTTCTTATTGCTGGGTGTTTCTCAACCATGGTTTCGCATTAAAATCAT

Fig. 9.101

Fig. 9.102

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AACAATGTGAATGTAGATACCAATATTGCAATTATGACTAATAATTGGAGGCAATTATTTATAGTAGTTATATATCAGT
ATATATATTTACCATTTATACTGATATTAATGACTGTGGCTTTTTAATAGTGGCCACAGAAGTCACATAGCATGGTTTA
GTGAAGTTTGGTTCACATTTGTAGGCACTTCAGAAATATCATCTTTGAGAACACACACACACTTAAGTTCAGTGAGAAC
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AAAATTATTCTTTACAATAATGAAAGACATTTCCAGATTCAAAAAAAGGGAAAAATAAAAACCATGGAAATAATATAT
TTGGAATTACAGGGCTACTAGCAATCTAAGTGTTGTGGAAAATCTTGGTAAATAGTTCAGTGAAATTATATAGATAGAA
GATTGATTGAATAAACCTACTCCAAAGCATTGATATGCCACAGCATTCTTCCTTTGGCTGTGTTCTGCCCAATATTTTA
ACAAGGGGTTGCATCAAAACAGAGTGATGCTGATCAACTCCTGAAAAATATTTAAAGTTAAAAGAAATGCTAAGCAAAA
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TGGTACAAGAACAAACACATAGACCAGAGAAAAAGAATAGAGAACCAGAAACAAGACTGCATACCCACAACCATCTGT
TCTTTGACAAACCTGATAAAAAACAAGCAATGGGGAAATCATTCCCTATTCAATAAATGGTGCTGGGACAACGGGGCTAGC
CATATGCAGAAAACCTGAAACTGGACCCCTTCTTACACCATATACAAAAATAAACTCAAGGTAGATTAAAGAATTCAAT
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TGACAAAAATGACAAAAGCAATTGCAAAAAAAGCAAAAAATTGACAAATGGGATCTAATTAAACTAAAGAGCTTCTGCA
CAGGAAAAGAAAATATTAACAGAGTAAACAGCCTACAGAATGGCAGAAAATTGTTGCAATCTATCCAGCTGACAAAGGT
CTAATATACAGCATTTATAAGGAACTTAAATAAATTTACAAGAAAAATACAACCCCATTA AAAAGTGGGCAAAATACAT
GAACAGACACTTCTCAAAAGAAGACATTCATGTGGCCAACAAACATATGAAAAAAAATCACTGATCTTTAGAGAAAC
CCAAATCAAAACCACAATGAGATACCATCTCACACCAGTCAGAATGGCTGTGATTAAGAAGTCAAAAAACAACAGATGC
TGCTTATGGAGAAAAACGAATGCTTTTACTCTGTTGGTGAAAGTGTAATTAGTTCAACTATTGTGGAAGACAGTGTGG
CAATTCCTTAAAGACCTAGAGGCAGAAATATCATTTAACCCAGAAGTCCCATTTACTGGGTATATACCCAAAGGAATATA
AATCATTCTGTCATAAAGACACATGCACGTGTATGTTCAATTGCAGCACTGTTTACAATAGCAAAGACATCTAAATGCCT
ATCAATGACAGATTTGGTAAAGAAAATGTGGTACATATACCCATGGAATTATGCAGCCATAAATAAAGAATGAGATCC
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CATGTTCTCACTTATAAGTGATGATCAGAACGCATGGACACATTGGGAGTGGGGAAACAATACACACTGGGGCCTTTCA
GAGGGTAGGAGGGTGAGAGGAGGGAGAGGATCAAGAGAATAGCCAATGGATGCTGGGCTTAATACCTGGGTAATGGGAT
GATCTATGCAGCAAACCACCATGGCACAATTTACCTATGTAACAAACCTGCACATCCCGCACATGTACCCATGAACCTTA
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CAGCCCCAATAACATAGGTGTTTTTGTTTAGCAAGGAATAAGAAAAGAAATGGATATTGGGTGGATACCTAATAGTAGTC
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GCACTTTGGTAGGCCAAGGAGGGTGGATCACTTGAGGTCAGGAGTTCGAGACTAGCCTGGCCAATATAGTGGAACCTTG
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TTCTACAAATGCATTTGCTCTGTCTGATAAAAATCATTTTAACTTTGCTTCTCCATCCAGCTACCTTCTTTTTTTTACCC
TTCTGTACTGCTTCTGGTATTCGATGGTTTTTCCCTTTCCCTTCCCCCATTTTTTGAACGATAGACTACTTTTTTTTCA
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GAAAAGAAAGACATCTGGCAATTAGATCTGACTTTATCCATTCTGGCTGTTATAACAAAATAGCATACACTCAGTAGCT
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GGTGAGGGTGTGCTTTCTGGTTCATAGTTGGCACCTTATAGTTGTGTCTACATGGTTAAAGGGGCGAAGGGTCTCTCTT
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AATATACATTTTGAAGGGACACAAACATTTAGAGCATTCCAAGGTCTTATTTGTTTTTTCAGTGGTTAAGAGTTTCGTCA
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CTTGTCATTATTTCTTAAGCAATACATTATAACAATTACTTACATAACATTTACGTTATACTAGGCATTACAACAAATC
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AGTGCAGCCTTAACTCGCAGTCTCAAGCGATCCTCCTGTCTCAGCCTCCCTAGTAGTTAGGACCAATGTGTGGGCCA
CCACCTGGATCATTTAAATTTTGTTTTTTGTGGAGATGGGGTCTCTACAAAAGAGATGTCACTTAGGTAGGTCTCAAAC
TCCTGGCCTCAAGTGATCCTCCTGCCTTGGCCTCCTAAAGTGCTGGGATTACAGGCAGGAGGTACCACACTCAGCCTGT
ATAATGTTTATAAGCACAAAATAAAATTTGTAGAATTAAAAATGAAACCAGTTGTATTAAAAACAATTATACAAATATTA
AAATAAAAATTTGGTATAGTTATATATGTGCATCTTTATTAATGTATTAAATCATAAGATCCAGCAGATCACATATCTA

Fig. 9.103

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ACATACTTAATTTTGAAGTGCTTGCAAGAAGTCTAATGAGATAAGAAGGTATCTATGATTTTACTGGCAACAAAGTCA
CAAACACAGTGGTTTGGTAACTATATTCATAATTGAAGAAAATGGTATTTTTCAGTTACAAGTTAGTAAAAATACAGA
TGTAACCTTGTATACAAGTTTACCAATCTCCTGAATTCCTTGTGGACTCCAGGTTAAAAACTACTAAATGGTAGAGTA
TATAGTTACAAGGAGTCAGATTTAAGCTCATTTTAAGTATATGTATTATGTGTATGTAAAGCCTTAATAATTAAGATGT
CCTAGAATGAAAGGATTTTCTCACAAAGTAGTGATCTACCTGTTAATGGAAACGTCCAGTTAGCATCTAGAAAAATATT
GAGGATGTTGTAGTATAACACTAATTAATATTAGCTAGCATCTACTGAGTGCTTGCTGTGTTTTATGCACTGGGTTAGG
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AGGAGGCAGAGTTTGAATGAGCCAAGATCACACCACTGCACTCCAGTCTGGACAACAGAACGAGACCCCGTCTAAAAA
AAATTAATAATAATAATAATAAAGTTGACGGCAATAAGTGGCAGAGTATGAACTCTAACCCTATCTAGGTGTCTCC
AAAGCCTATAATTGGAGAATATTTTGATAATAATGTAGGAGAGAGATTGGTGAGAGAATTAGAGATCACCTTGTTTCATC
CTCTTTATTTGATAGATATGAGGACACTGAGAACTCAAAGAAGTTAGGTGACTTACTCCAGGTTACACAGTTTATAGCA
GAGCCAGAAATTGGACTTTGATGCCTTTTTTATGTGGAACATGAGCTTTTATTATTAGCTCTTCATCTGGTGGAAGTG
GAACACAACCTGAAGAGAGAGGCAATGGACTACACTATGGTTTGGAAACAGAGTGTATAGTAATTTCTTATTTTCAATTTAG
ACAACAGGGATATGCCTGAAAGTGCTTTTACCCATGTCATGCATTTATTTCACAATGAACACAAAATTTACTTGAGTAAT
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CGCTCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCTACTTGAGTAATATTTTAAATGTAACCATAGTGAACCTGT
CCAATACTAAAATTTGTGCCTTTGATAATATTTATATTATGAATAAAAAATATGCTCTTTTAACCATGTCCTCATCTATT
TTACCAGAAAGTATCTTGTGTATTGTCAATACAAAGTATCTAATGCATGAATGACTGAGTATGATTGTCTGGTTTTCTT
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GTTCAAGATGTTGACAGAACACATGGTTTTTGTGTTTCTTACTTTTCTTAATACCCATTAAAATAGTAGAAAACTAA
ATCTGTATAATGCACAACAATAAAGAGAATGGGGAAAAGGCCATCAGTGGTAAGAGATTTTATCGAATTTCTGGAAGAT
GAAATATGAGTGAGTGGTGTGATTACTGAAGCAAAACAAAGTCAATTATTGCAGAGAATATTTGTAGAGGGAGCTACA
CCCAAGAAGAAGCCCTAATAACATGGCAGAAGCTAACAAGGCTCCAGTTTCAGATAAGCCAATGCTGATGGACATTGAA
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TTCCTGTGTCTTTGCTCAACCAATCAGTTGATCAGCTCTGCTTCTGCAGGTTGGTTGGGTGTCAGTTAGCAAGATAGGG
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ATCCTTCTCTGAGTGGCCTACTAATCTGACATAGGAATCAGAGAAAAGGAAATGTAATCTTAACCTTTCACTTTGCA
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CAGACATATCTTTTACATCTTTGAAATTTAAAGCTCAATGTGATGGTCTGTGTCTTGACTCTATTTAGCCAAACCACTG
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CAGTGACAGTGCAAGTCATCACCTTTTACTTTTTTGCAATACCAAGCTGTTTTTAAAGAGTCTCTTGCCATCCTGCC
TATGACTTTGTGCAAGTGAATCAGTTGAGTTCATGGGAAAGCAAATAAAAGGATGGTCTCTGTTTATAGACATAGCCCT
GTCTTAGAAATCTAATCTCTCTTATCACTCTCATCTACAAAGACTGTCAAGGAAATGTGTCTCTCTGCCCATGGAGA
CAGATTGGGCATCTCACATGAAAAAATATGTCTTTGAACTTGCCCTGAGAAATTCAGTAACTTTTTCTCACCAGGAT
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AGTATCAGAAATTTATTTCTCACAGTTCTGGAAGCTGGGAAGGCCAAGGTCAAAGCACCAGCAAATTTGGTGTCTGATG
AGGGCCCACTTTCTGGCTTATAGATGGTGCATTCTACCTGTGTCTCACATGGTGGAAAGGAGACAAGTCAGGTCTCTGG
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TAATATCATCACATTGGTGATTAGGTTTCAGCAGATGAGTTTTGGGGGAATACATTCAAGGCTGCAGCAAGGTCAAAAAG
AATATTGCATCATTTTGCCTTAGAGACCTTTCTGTTCTAGCTACATTTTGTATTATCTATATGACACAATAAAAAAGAAA
GTCAGGAAACCTAAGATCTTATACTTCAATAGAGTTTTTATAGGACAACATTGATTAATGGCTACAGTTAATACAAAAA
CTCTAACAGCAGCAATCAAATATTTCCATTCATCACACTTGAAACTTGCCCTCAGGTCTATGGTATTTTGAAATTTTTT
GTTGTTGTTTTGAGAGTGACATGAGTGCAGGCCAGCTTCATGGATATACAACTATACATTACATGGGTTCACACC
TAGATGGGCTCTTACATCATGTAGCTGGTCTTACCTGGGAGAGAGCTAAAAATTAAGTTGCAGTCATCAAATTACTGAT
CAGATACTGAAATTAACCTGAAAGATGTTGAGAACCAGAGAAATTTCTCAATCATGTTAATAATACTTTTCATGCATTT
GACATTTCTTTTTTCTGCTCTTCTCTCTACCTGTAGAATTTTCAAGTTCATTATTAATACCATTTGTTGGCCCTTTG
TTCTCTCTCCAAAGACTTAAAGAAGAAGTTATGATTAAGCATATGAAATATGCATACCAGGTTTTCTATAAAGATTCC

Fig. 9.104

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TCGGCATCTCTGCCAAGCTGTCATATGAGCACTTGCATGTTGTTATATAAAAACACATATTTTAAATAAATTTTAGTGCT
TGTTCTAAGCAAAAAGCACATATTTTAAAAATCTGGAGTTTTCTGTTTCAATTTGAGAGAACTACATTTTGCTTAATTA
CTATTTAAATCACTGTAACACAATATTGATGACAAAATTTGAGTTTGCTTTTTTAATAAACTGTCAATCAAAGAAATAAC
ATAGAACTTTGTTTTTCTCCTCTGTTAAGGCATGGCTTAAGTGTAAATTTACTTGAAAACATTATACTGTTTGACAGGAA
AACAGAACCACCTTGAATTATTCTAACCCATTGAGATTTAAGAAAATGCATCGTAGCAAATTAAGTGTATTTGGTTTT
GATCATTTCTGTCTGTATCATAGAAATATGGTTCAGTAGTGAATCAATGAGAGATAAAAGGTGATCATGGTCTGTGAGAT
AAATGCTGCCCCCTCTGAGTTGCTTCCAACGTAGCCAGCTTCAGTGTATCAGTTAGAGTAGTTAATACTAAGTGTGTAA
CAGATAAATACCCAAATCCCAGTTGCTCAAACAATTGAAGTTTATCTCTCATTCATGCTAAGTCAGTTGATGTTTCTGG
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TCTCCCTGCTACCATGTCTCACAGGACCTCTTCTTGGTTCAGTATTGCAACTATCTCCAGTTTTCTCTTTTCATCT
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TTTTATTGCCTTACACTATACTTCCCATGCCTCTGATTTCAATGATCACTTATATACTAAGGACTCGCCAATTTGTATA
TTCAGCCCATATTTGTCTCTGGAACCTCAGACCTGTATATCCTACCGCTTTTTCAATATCTTCCATTGAGTGTTCAGA
GGCATTTTAGATCCAAAATATCCAAAACCTGAAATTTCTGTTTTCCCTCACAGCCTGCTTACTGCGACCTGATTTTTCTC
TTGTGGCCCTTAAACTATGAATATCAACAGTCATCCAATTTCTTACCTTCATCTTCTCCCTCAGTTGCTGTAAGTAC
CCTATCACTAAGGCCTGTGATTTTACCTTCCAGTTATGCGCTGCATCTATCCCTCTCTCTTCATCTTTCCCGGGGCCG
GCCTGTCTGTGCTGGCATTACCTGTTTCTGTAACACAGTGATAGCTGCCTTGATCCATTTCTTGGCATCTGCTCTTGC
CCCTCATCCAATCTGTTCTCCATATTGCAGTCAGTCACCTGTTTGAAACAGAAGTCCAACCTTATCACTCACACACAC
CCTTGCCTTAAACAGTGTAGCGGTTTCCCATTTGTTCTTAAGACTATTAACGTCATTTTTGTGTGATTGTCCAGGGGCT
CAATAATCTGAATCACACCTTGTCTCCAGCCTCCTTTCTGCCTTATCCTGAAGTATCCCTGCCTGTTTCACTCCCTCA
GCTTCTTGTACACCGACTTTCCACCCAGTGCCTCAGGGCACTTTCTCATGCTGTTCTCTCTGCCAGGAACACATGCTTC
TGCTTTTCTCTCCCTTTACCTAGTGAATCCCTCCTCATCCTTCCCATCTTTGTAAATTCATTTCACTTCAGTTTAGAAGACTT
CTCTGGACTGTTATATTGTGAGAACTATCTTCTATATATGCTGCAGAAGTATACATACGCTGCATACTTCTTTCACTGT
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AGAGTAAGTCAACAACCTTTTTTTTTTTTTTTAGCAAACTCTAAATGTCTGGATGATCTGATAGATGCAATTCATATT
CGAGTCAAATGTTGAATATGTATTACAATTTAGGCTGTTCTTGAAGCTTTATTAAATTGAGCACCAAATCTTCTGAAGC
TAATGGAGGTAAATCATAGTCAAACGCCCTTTTAACTCAAAGAACCATGCCCCCACTATTTCTCTTTCTCAAACTA
TTGCTTCTTTCAGGCTAGGATATTTGGGATATTAAAGGGCTCTTTTAACTCAAATAGCAAACCATAGCCCTCATCTT
CTTATTTAGGAAGACGGTCTTTAACATTTAATTCTGAGCACTTACCTTTTCCGTAAATGAAATTCGTCTTCTCATTTGG
ATATTGGCAGAGAGCCAAGATAAGGTGAGCATTGAGCAGGATAAACTAGATGATTCCTTAATTCAGCAAATATATTTTG
ATTCCAATTGTGTACCAGAAATCTTCTAGGTACTGGGAATTTATCAGTGAACAAATAACATTTCTATCCTCACAGAGC
TTACATTCTGAAAGATAGGTAAATAAATAATTATCTATCTATCTAATTGAATCACTCTTCTCTCTCTCTCTCTGTCTT
TCTCTCTTTGTCTCCCCGCTTGGGTCTTTCAAAGACATTTCTGCAGGAGTACAGCTATGAGCTGTTTAGCAGTCAGTAC
AACCTCTGGAAGATGGGTGTACCTGCCCAATAAAGGGGATCTGAGAAGACCGTCAGTAGCATCTACTATAGAAGATAAA
TGCAGTATTTCCAGGCAGGAGATTATGATGGCTTGGACTAGGGTGGTAGCTGTGTTTACGGTGAGAAAGATAAGATCTGG
ATATATTTCTAAGGTAGATTTTACAAGTTTGTCTGCTTGATTGAGATCAGTTGCAAGGAAAAGGAAAGTCAAAAATGACT
ATATAAATTTCTGGTCTGAGAACTGTGCAAAATAGAGATACTGTTTACTGAAATGGAAAGGGAGAGATCAAGATTTTCAT
GTTAAATTTGGTACATTCATTGAGATATATCAGTGGAAAGATAGGCAGTTGGTTATGGTAGTCTTGAGTTTCAAGTTGGT
TTTACAAATCTAATTTGGGTATTGATGATTTTAAACTGAGGAGTTACCATAGATAAATAGGAAAAGCCACTCTAATGT
TTTGAAGTTGGAAATTCAAAAGGAAATGGAGCTTTTGAAGAAGGAGCAGCTGGTGAGGCAGAAATACAACCTAGAGTGAG
GAAAGGCTTGGAAAGTCAATGAAGATAGTGTTTTACAAAGGAATGCATGATTAATTGTGAATGCTGCTGATTACTTAAGT
GAGAAGTGAATGACAAATGAATTTAGCACTGAGAAGACCTTGGATGATCTTGAGAAGAGCTGTTTCAAGTGGAGTAC
TATAAACAAAAACATAGGAGTACATTCAAGAAGAAATGGTGGGAGGGCAAGAACTGAATATCATGAGTTTGGAAAACCTC
TTTTGAGGAGTTTTACTTTAAAGAATTTAGGCAGAGAAATAAGATATATAGTAGTTAAGACTTAAATTGGACATATTAT
GGCAGGTTGCTATGTAAATGGGAAGAATCCAGAGAAGAGGAGGAAATTAGTGATGCAACAGAGACAGTAATAACTGGAG
TGATATACTTGAAGAGGAGAAAAGAGATGATATCAGGCACATAAATGGAGGATTTGATAAATGTGATGGGGGCTGCAGA
GAAAATTTGTTTTCTGATTTGCTATTATTTTTTTTCAAAGAAATAGGCTATCAACTGAGTGTGAGGATAGAGCAGAAGCT

Fig. 9.105

GTTAAAAATTTAAGGAGAGTTGTGAAATAATCATCTGTAAGCAGGAGTGTGAATGAAGTAGGAAGATGTAGTAATATTT
CTGGACAGCCCTGAGAGCTGAAATATTGTAAGAAATAATCTGAGGAAGTTCTGAGTACTTAGACAGTTGAGGTTAGGGG
GAGCATTTTATTAAAAACAATCACAAAGAACATGGGAAAAACAGATACCAACTTGGCAATTGTCATAGTATTAAATAAAA
CTAATTTTGGAAAATTAACTAAAAAGATGTTATATGCATTATGATGTTTCAAAATAGACTAGGGCCAATCAGACCCAA
TAGTTTCTTTTATATAAGACCCCATAGTTTAATTATATAATTAAGTGTACGTTCAAATTTTGATAAATGATAAATATAT
TGTACCACCATGGGAATTTGAAAATGCGACCATTGTACTGGGATATTATCATGTAATATTTTAATTTTACCTCTTAAT
TATCAAAGTGAAAGATTTCCATGCAGTATTATTTGGTTGTGGGGTAATTGTTTAATTTATTTGGAATTTTATAGTGCTT
TTGCTTAGCCACATTTTATTTTGTGGTCAAATTCAAATGCTGATCTAATGGCTTCAGAGTAAAATGAGAAAGGTCAAAT
GGGACTGGAAATTAGGGTTTTCTTCCTGTCATTAGGTATATTTATTTATAAAATAACTGGAAAATTCCTCATGTATTTA
ACATCTGTTCTTCTTTCCCTGGTCACTGCATATCAAGAAGACTTTGACATTGAGGTTTGATACGGTTTGGCTCTGTGT
CCCCACCCAAATCTCATCTTGAATTGTACTCCATAATTCCCAAATGTTGTGGGAGGGACTCGGTGGGAGATAATTTGA
ATCATGGGGTAAACTTTCCCCCATACTGCTCTCATGGTAGTGAATAAGTGTCAAGATCTGATGGTTTTATCAGGGGT
TTCTGCTTTTGCATCTTCTCATTTTCTCTTGCCACTGACGTGTAAGAAGTACCTTTTGCCTCCTGCCATAATTGTGAG
GCCTTCCCAGCCATGTGGAAGTGTAAAGTCCAATTAAACCACTTTTTCTTCCCAGTCTCGGGTATGTCTTTATAAGCAAT
GTGAAAATGGACTAATACAGTAAATTGGTACCAATAGAGTGGAGTGTGATGAAAAGATACCTGAAATGTGGAAGCGAC
TTTGGAAATTGCCAACAGGCAGAGGTTGAAACAGTTTGGAGAGCTCAGAAGAAGACAGAATAATGTGGGAAAGTATGGA
GCTTCTTAGAGACTTGTTAAATGGCTTTGACCCAAAGCCTGCTAGCAATATGGACAATAAGATCCAAGCTGAGGTGCTC
TCAGATGGAGATAAGGAACTTGTTGGGAAGTGTAGCAAAGGTGATTCTTATTATGTTTTAGCAAAGAGACTCACAGCAT
TTTGCCATGCCCTAGAAATTTGTGGAACCTTGAACCTGAGAGATGATTTAGGGTATCTGGTGGAAAGAAATTTCTAAGCA
GCAAAGTATTCAAGAGGTGACTTGAGTGCTGTTAAAGGCCTCAGTTTTTATAAGAGAAGCAGAGCAGAAAAGTTTAAAA
AATTTTGTAGCCTGACAATGTGATAGAAAAGAAAAACCTATTTTCTGAGGAGAAATTGAAGCTGGCTGCAGAAATTTGC
ATAAGTAACGAGAGGCCGAATGTTAAGCCTTCAAGACGATGAGGAAAATGTCTCCAGAGTATCTCAGAGGTCTTCACAG
CAGCCCCTCCCATCACAGGCCTGGAAGCCTAGGAGAAAATGGTTTTGTGGGCCAGGCCCAGGGTCCCCGTGCTGTGTGC
AGTCTAGTGACCAGGTGCCCTGCATCCAGCCACTCCAGCTGTGACTAAAAGGGGCCAAAGTACAGCTCGTGCTATGGC
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GTTTCGGGAACCTCCGCCTATATTTCAGAAGATGTATGGAATGCCTGGATGCCCAGGCAGAAAGTTTGCTGCAGGGGCAG
GGCCCTCATGGAACCTCTGCTAGGGAAGTGTGGAAGGGAAATGTGGGGTTGGAGCCTCCACACAGAGTCCCTACTGG
GGCACTGCCCTAGTGAGCTATGAGAAGAGGGGCCACAGCCTTCAGACCCAGAAATGGTAGATCCAATGACAGCTTGAAGC
ATGTGCCTGGAAAAGCCACAGATACTCAACGCCAGCCCATGAAAGCAGCCAGTGGGAGGCTGCACCCTGCAAAACCCAA
GCAGCAGAGGTGCCAAGACCATGGGAACCCACCTCTTGCATCAATGTGACCTGGATGTGAGACATGGAGTCAAAGGAG
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AATTTCTCCATTTGGAATGGCTGCATTTACCCACTACCTGTACCTGCATTGTCTCTAGGAAGTAAGTCTTGCTTTT
GATTTTACGGGCTCATAGGTGGAAGGAACTTGCCTTGTCTCAGATGAGACTTTGGACTGTTGACTTTTGGGTTAATGCT
GAAATGAGTTAAGACTTTTCAGGGACTATTGGGAAGGCATGATTGGTTTTGAAATGTGAGGACGTGAGATTTGGAGGGCC
CAGGGGGGAATTATATGTTTTAGCTCCGTGTCCCCACCCAAATCTCATCTTGAATTGTACTCCCATAAATTTCCACATGT
TGTGGGAGGCATTTGGTGGGAGATAATTAGAATCATGGGGCAGTTTCTCCACACTGTTCTCGTGGTCTGTAATAAGTC
TCACAAGATCTGATGGTTTTATCAGGGGTTTCCACTTTTGATCTTCTCTCATTTTCTCTTGCCACCAACTTGTAAAGAG
TGCTTTTGCCTCCTGCCATGATTGTGAGGCCTCCCCAGACACATGGAAGTGTAAAGTCCAATTAAACCTCTTTTTCTTC
CCAGTCTTGGGTATGTCTTTATCGCCACATGAAAACAGACTAATAACAAGTTATTCTATGAGTTAGAAATAATTCCTCT
AAAAGTAACACTTGCTGAGAATTTCCCTACCTTTTTCTGGGCTTTTAAAAATGCATCTTATTCCTCATCCCCCTAAAGTGG
GTGTGTTAGTCAGGGTTCTCTAGAGGGACAAAATAAGGAGATATATATATATATATCTCCTATATATCCTATATAT
ATATAGTATTAACCTCACATGATCACATGGTCCCACAATAGGCCTTCTGCAAGCTGAGGAACAAGGAGAGCCATTCCGAG
TCCCAAACTGAACTTGGAGTCCAATTTTCAAGGGCAGGAAGCATCCAGCATGGGAGAAAGATGTAGAGTGGGAGTCTA
GGCCAGTCTCATGTTTTACATTTCTTCTGCCTGCTTTATATTCTAGCTGTGCTGGTAGCTGATTAGATAGTGCCCACTC
AGATTAAGGGTGGGTCTGCCTTTCCAGCCCACTGACTCAAATGTTAATCTCCTTTGGCAACACCCTCACAGACACACC
CAGGATCAATACTTTGCATCCTTCAATCTAATCAAGTTGACACTCAGTGTTAACTATCGCAGTGGGTAAAAGCCATTAC
TAGGCAACCCAAACCGTTGCTGCAGGTTGATGCAGCAATGAGTAGGAAGATGTGGTGGGAAGATGTGGGAGTATAAACTT
CGCCAGGCAGTAATATAGGTATATAATACATGAGCAGGGCTGCAATACTCTCTTAGGAACTGGACAGCTTTATGGGCCA
AACCAACCTCAAAACCCAGCTGCCCACCTGACACAAGACTGTCCATGGGAGTGAAGTTCTGCAGGGTTCTTTCCCATC
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CTACACAGCTCTTGTCTTGAAGAGCTGTCTTGCATTGTCAATACCTTGGTGCAACATATGTGGCTTCTGCCCCTTCT
GCCAAGTGGACCTCCTCACTTTGAGCACATGGTAAGCATCTCCATAGGTACACAAAGTTTGTAGCAACTAACTCCTAT
CTGGAGTTACTCAGACCCACTACCTCTCCTGTTAGTGATCTATGCTCTTCAAAAGTACAGTAAGCTCTGCTTCACTT
GTCTTGAAAACATGAATTTGTTCTAATGCAATGGACATATTGGAGAAGAATTTGATGAACAGCTAAAAAAGCAATAGT
ATAAATAATTTTCATGGCTCTACATGTAACCTCTGGAATGCATAACAATGCTTTTTTCCATAAGAAGATGTTTTTCTGT
GCTTGTAGATTATTTCTTTTAGACCTGTAAAGAATGCTTGGAAATAATTGATGAAGTGCCAAGTCCACTAAGTTTCTGTT
TTTAGCTTCTATCCTGAAGTTTCTCATATTCTTTACCTTTACATCTCTCAGTGTGAGCTCATATGAATCTGTGGCTTCA
GGTCCCATGTATATACTGATGACACCCAAATCTCTTCTTGAATGACCAAAGACCAAACCTTTCTGAGTCACAGATTTTAA
TTTACATCTGAATGTACCCCTAGTATTTTAAATTTCTTAAATATAAAAGTTTAAATATTTCTCATACCCACTTATGTATA
TGCATGTGTGAGTGTGAGTATGAGTGTGTGTACCACATATACTTACTATGTGTGTAGGCATTGTCCTATATGTTTCAT
ATCTATTAACCTCATTAATTTCCCAACAGCCTTATGAGTTAAGTCTTCTCATTTATGTCCTTATAATAAATGAAGAACT

Fig. 9.106

Fig. 9.107

TATTGTTTCTCATCCACTGTGACATTCTATGTTATTTTCAGTAAGATTTAACAAGCAAAGTGATGATGGATGAGAGTTCC
TTTGCAGCCTGGTGAGATTTTTACTTTCATGTAAAGCAAAGCATTTTGGGGCAACATCAGGGATAAACTTGTGGAATGAA
TCATTTGGTTTTTCTTTAGTATATCCTGTCCACCATTGGGAACCAGATTTCAGTAATTATGTTCTCTGGTTTTGTACTCAT
CATAATGTCTTGACCTCCAGCTCTGAAACAACATATTTCTGAGAGATGCTGATTGTATTTTGTGAGGTTCTGCAGGGT
CAAGTAAATTGAGGATGTCTTTATTCTGCATTTGACTAATCTGAATTAGATAGTCTTAGCCATTAAAACTTTAGAAAGC
ACATAATGACCAATTACAAGTGAGAGAATTTTTTTCAATAATGCTTTATATTTTCTAAGTACCGATACAGATGTTTTAT
TACATTTAAAACATGATTGCCCTACCACTTCACTTATATTTTAAAAACTAAGTTGCCATCTTGTAATAATAATACATG
GATATCATTTACAAGTGATATCAGTTAATACCTAATGTGAACATCCATTGTGTACTGAGTCTATGAAATACTACTAGTT
ACTATGAAAATGTGAGATTACCCGAATCGGGTGGAGCATCAACTTTATTATCCACATAGGTTACCCATTTCTTATATCA
TAGCTTTGGTTGTATAATTTATTTCAACAGAGTTTAGTTAAACATTTTCCACTGTTATTAGGTACTGCCACTCTACATT
CCAGTAAATATTGCCACCATGTTGCAGAACAAATTTTGGACAACCAGGCATATCTTTGGAGTTGAACTATGTGGAGGTA
CTACTGGTGCTGTGAGGGATCACCTGTAAACCTGGCAGTTTLAGGTGGCCTCTAGACGTTATAGGGTCAAGGGAGAAAT
TTTTTTTGTCTTGTGTTTAATACTGTGAGTACATTTAAATGCTCATGTAAAGAGCCAACTTAGAGGGAGTGGTTGA
AGACACAAGAGAGATAAGAGCTGATTGTTATGGCAAGGTCTCTGGCTAGGCAGAAGTGAATGGGTGGAAGAGAACAGAT
TTAGGAACTGACCTTCTATCAGGAGAACAATATTTCTTCCAATGTCATGGGGGAACATAGAGACTCAGGGGGTCTGTAA
CCATGTATGGCCCCCAAAGGACTGGACACATGGTGAGCAACTATGGTAGGATGAAGGGAGGCATGTTAGGAAATGTCCA
GCTCCGAGGCCTCCACGTGACCCCTAGGTGAGAAGCTGCCATGATTCCTTAATAGGGAGCAACACAAGACTAAAATGA
TGGCAATGTAAGGGTTAAGGAAAAGGTGATTTTAAAAGCTTCTTGAGGGTAAAGTCCATCTGTGTTCACTTTTGTGATC
TTTGTTCATCCTTCTCAATCTTGCATATTGTTGATCTCAAATAAGTTTATGGTATTTATTATGTTCAAAAAAGTTACA
GTTTCTTAAGTTTCTTCTATTTTCTTAAGGAAGGAGGTGATAGGATGGAGGGGGCATGATGCCAACTTGACTCTCCTGC
TTAGGACTTGCCCTTCTTTTAAATGGTGAACACAGAGTTACATTATTTAATGCATAATGTGAAAGTGACAGAGGCCCTG
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TAAGCCCATGGAACATCCTCTTCAAATGAAGTTGACGTACGTCTCATCTCCCTGGATTGGATGCCCAACCCTCACCTT
GAAAGCAGCTTGTGGATAACTCAGGGCAAACAGAAGCAGAACCCTCATGAACCTTCTCTCTTCTGGAAATTTTGGATGG
CGTTACAGAATGAGGCAAAAATAAAGTTACCTTCTTCTCTTTTCTTGGATTAAAGATCCCAGAGCCATTCTCCTCTGTG
TTCTCTACTTGGACCTTGTATAGTGTAGGATATAGCCAGGAACCTAAGGATCCTCGGTTGACTCATAGTCGGCAGAGGGT
GTTATGGCCCTCCTGAGAGAGTGACCTTGAGGAGAGAATCTATTCAGATGGTTTTATGACAAGACCAAAGTAACATTTT
TTAGTTACCTCTTAGCCAGAGTTTAGTGTTAAGTTTGCCCTGAAGTATAGTTTTTCAAATAATGAGATCAGGCATATGC
TATTTTGACTAAGTAGAATACCCAATAAAATAAGTCATTCAAGTTGTGACTGAAAGTTCCGGATGGTCTTATGGAATCA
AGGTGACCAAATAACATGATACTAAACCAGAAGTGAGTCATGTTGTTTATCATACTGTTTTTATAAATTTTGTATAATA
CAACCCCATACGACCCACACAGTTCAAACAGTAGTGGTTACTCACTCCATGGAGAAGATGTCAAAGTAAAAGGAAT
AATACATTGGTTATGATGTCTGAAATTGGGCACATAAGCATTTAGTTTTGTTGGTGAATAATGTTTCAAGATGAAAATTG
ACACACTCTTTGGATTATCAAGCTTTTTTAAAAGTATAATTAATCAGTTTTTTAATGAAATATACAGAAATATAATGGC
AGCTCTTCTATTAGTGAAGTTTCCTGAGGCAGCTGGATTGATTGAAGGCTTTACCTGGTTTCATTTGGATTCTACTTGTA
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TATCAAACATTAAGATGAGATAAAAATAGCTTTTAAATCAAGTAAGTTAAATTAATAACTGTTAAGGTTCAAATCATG
AATTAAATCTAGATGTTTGGCAAAGACAAAATCTTTATAAAAGGATAAAATAGTCTGGGAATAAAGTATTTGTCTATTC
CCATATCAAATGAGAACAAAACCTAATAATTTTATTAACCTTTAGGTTTTTTATTGTATTAAATAACATCATTTTTTATTAC
TTTATCACAAGGTATATTTTTCTTTATTATATGTCTTTGTAGAAAACCTAGTAACATTCCTGTGTGACTGAATAGATTA
ATCACAGATATACCCATCTGGAAAAATGCATGCTACTATAAGAGATGAGTGAAATATATAAAATTTATATTTTAAATTCT
TATGCTAATTAGTTAAATGGGGGAAGATGTATGTCCAGAATATTTGCTCTTAATAGGACATCGTAGTGAAAACCATTC
AATGATGATGACTAATAAATGTGTATCTTCAACATTGTATAATGCCCAGGAATATTTCCAAATAAAGAATTTCTAGGTA
AGATGTTTTAACAAATATATGATACTTTTGTCTTCTGGGTGAATGTTCAAACCTAATTTCTGGGGATCATTGTGCTCAGT
ACTTATAGTATCATTAATATTATTGTTGACATATAATAATAACCAAAAATATCCTGGTCAGATAGGTTTTTTCTTTTTT
AGCTGATTCATAATCATTGTACATATTTATGGGGTACAGAATGATATTTCAATATGCGTATACAATGTGTAATGATCAA
ATCAGGGTAATTAGCGTATACATCACCTCAAACATTTGTCTATTTCTTTGTGTTTTGAACATTGAAAATCCTCTCCTCTA
GTTTTTGAAAATACACAATGGATTATAGCTAACCATGTTTACCCAACAGAGCTACAGAACACCAGAACTCATTCCTCTC
ATCTAGCTATAATTTTATATCTGTTAACCACCTCCTTCCCATCCTCCTCTTCCCCATCCTTCCCAATCTCTAATACCC
ACAATTCTACTTTACTACCATGAGCTGAATTTTTTTATTTTAGCTCCCTCTTATGAGTGAGAATGTGCGGTATTTATCTT
TCTGTGCCTGACTTATTTCACTTAACATAATATCCTCCAGGTTTCATCCACATTGCTGAGAATGACAGGATTTTCATTATC
TTTTTGTGGCTGAATAGTATTTTCATTGTGTGTATATAACCACATTTTCTTTATCTATTTGTCTGTTGGTAGACATTAAGG
TTGATTCCATATCTTAGCTGTTATAAGTAGTGCCGCAATAAACATGATGGTAGAGGTATCCCTCTGATATAATTGGTTTC
CTTTCCTTTGGATAGATAACCCAGTAATGGGATTGCTACATCATATGTTAGTTCTATTTTTTAGTTTTTAAAGAAATTTCC
AGATTGTTTTCCATAATGGCTATACTAATTTACATTTGCACCAACAATGTATAAAAGTTGCCTTTTCTCTGCATCTTTG
CCATCATTTGTTATTTTTTCTTTTTTCAATAATGGCCATTCTAAGTGGGGTGAGATAATATCTCATTTGTGGTTTTGATGAT
TAGTGATGTTTCAGCATTTTCTCATATACCTGTTGGCCATTTGCAAGTCTTTTGAAAAGTGCTATTTCAGATATTTTGCC
CACTTTTGAATCACCTTATTTGTTTTTTTTCTGTTGAATTGTTTGAGTTCTTGTATATTCTGGATATTAGTCCATTGTC
AGGTGAATAGTTTGCAAAATATTTTCTCCCATTTCTACAGGTTGTCTCTTCACACTGTTGATTGTTTTCTTTCTGTACAG
AACTTTTTTAGTTTTAATATAGTCCCATTTGACTATTTTCGTTTTTGTGACTGTGCCTCTGAAGTCTTAGCCAAATAGT
TTTTATTTCTTCAAATAGCTTTGGCATTGAGTACCACTTCGCTCATTCAGAAAAGCAACCACCCCAACCAAAAGAAT
AGGTTTTTAATTTTTTTTTTAATTATTATTATACTTTAAGTTTTAGGGTACATGTGCACAATGTGCAGGTTAGTTACATA

Fig. 9.108

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TGTATACATGTGCCATGCTGGTGTGCTGCACCCATTAACATCATCATTTAGCATTAGGTATATCTCCTAATGCTATCCCT
CCCCCGTCCCCTCACCTCACAAACAGTCCCCAGAGTGTGATGTTCCCCTTCCTGTGTCCATGTGTTCTCATTGTTTCAGTT
CCCACCTATGAGTGAGAATATGCGGTGTTTGGTTTTTTGTCTTGGCAGATAGTTTACTGAGAATGATGATTTCCAATTTT
ATCCTTGTCCCTACAAAGGACGTGAACTCATCATTTTTTATGTCTGCTTATTATTCCATGGTGTATATGTGCCACATTT
TCTTAATCCAGTCTATCATTGTTGGACATTTGGGGTGGTTCCAAGTCTTTGCTATTTTGAATAGTGCCACAATAAACAT
ACGTGTGCATGTGTCTTTATAGCAGCATGATTTATAGTCCTTTGGGTATATACCCAGTAATGGGATGGCTGGGTCAAAT
GGTATTTCTAGTTCTAGATCCCTGAGGAATCGCCACACTGACTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAAC
AGTGTAAGAGTGTTCCTATTTCTCCACATCCTCTCCAGCACCTGTTGTTTCTGACTTTTTTAATGATTGCCTTTCTAAC
TGGTGTGAGATGGTATCTCATTGTGGTTTTGATTTGCATTCCTCTGATGGCCAGTGATGGTGAGCATTTTTTTCATGTGT
TTTTTGGCTGCATAAATGTCTTCTTTTGAGAAGTGTCTGTTTCATGGCCTTCGCCCACTTTTTGATGGGGTGTGTTTGT
TTTTCTTGTAATTTGTTTGAGTTTCAATGTAGATTCTGGATATTAGCCCTTTCTCAGATGAGTAGGTTGTGAAAATTTT
CTCCCATGTTGTAGGTTGCCGTGTTCACTCTGATGGTAGTTTCTTTTGCTGTGCAGAAGCTCTTTAGTTTAATTAGATCC
TATTTGTCAATTTTGGCTTTGGTTGCCATTGCTTTTGATGGTTTAGACATGAAGTCCTTGCCCATGCCTATGTCCTGAA
TGGTAATGCCTAGGTTTTCTTCTAGGGTTTTTATGGTTTTAGGTCTAACATTTAAGTCTTTAATCCATCTTGAATTGAT
TTTTGTATAAGGTGTAAGGAAGGGATCCAGTTTCAGCTTTCTACATAGGGCTAGCCAGTTTTTCCAGCACCATTTATTA
AATAGGGAATCGTTTCCCCATTGCTTGTTTTTCTCAGGTTTGTCAAAGATCAGATAGTTGTAGATATGTGGCATTATTT
CTGAGGGCTCTGTTCTGTTCCATTGATCTATATCTCTGTTTTGGTACCAGTACCATGCTGTTTTGGTTACTGTAGCCTT
GTAGTATAGTTTGAAGTCAGGTAGCGTGATGCCTCCAGCTTTGTTCTTTTGGCTTAGGATTCATTTGGCAACGCGAGCT
CTTTTTTGGTTCCATATGAACTTTACAGTAGTTTTTCCAATTCGTGAAGAAAGTCATTGGTAGCTTGATGGGGATGG
CATTGAATCTTTAAATTACCTTGGGCAGTATGGCCATTTTCACGATATTGATTCTTCCGACCCATGAACATGGAATGTT
CTTCCATTTGTTTGTATCCTCTTTTATTTCTTGGAGCAGTGGTTTGTAGTTCTCCTTGAAGAGGTCCTTCACATCCCTT
GTAAGTTGGATTCCTAGGTATTTTATTTCTCTTTGAAGCAATTGTGAATGGGAGTTCACATCATGATTTGGCTCTCTGTTT
GTCTGTTGTTGCTGTATAAGAATGCTTGTGATTTTTGTACATTGATTTTGTATCCTGAGACTTTGCTGAAGTTGCTTAT
CAGCTTAAGGAGATTTTGGGCTGAGACGATGGGGTTTTCTAGATATACAATCATGTCGTCTGCAAACAGGGACAATTTG
ACTTCTCTTTTTCTAACTGAATACCCTTTGTTTCTCTCTGCTTAATTGCCCTGGTCAGAACTTCCAACACTATGT
TGACTAGGAGTGGTGAGAGAAGGCATCCCTGTCTTGTGCCAGTTTTCAAACGGAATGATTCTAGTTTTTGGCCATTGAG
TATGATATTGGCTGTGGGTTTGTACATAGGTCTTCTTATTTTGAAGATACGTCCCATCAATACCTAATTTATTGAGA
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CGGTTCTGTTTATATGCTTGATTACATTTATTGATTTGTGTATATTGAACCAGCCTTGATCCCAGGGATGAAGCCAC
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CATCAAATGAGTTAGGGAGGATTCCCTCTTTTTCTATTGATTGGAATAGTTTCAGAAGGAATGGTAGCAGCTCCTCCTT
GTACCTCTGGTAGAATTCGGCTGTGAATCCATCTGGTCCCTGGACTCTTTTTTGGTTGGTAAGCTATTGATTATTGCCACA
ATTTTCAGATCCTGTTATTGGTCTATTTCAGAGATTCAACTTCTTCCCGGTTTAGTCTTGGGAGAGTGTGTGTGTCCAGGA
ATTTATCCATTTCTTCTAGATTTTCTAGTTTATTTGCGTAGAGGTTTTTGTAGTATTTCTCTAATGGTAGTTTGTATTTT
TGTGGGATCATTGGTGATATTCCCTTTATCATTTTTTATTGCATCTATTGATTCTTCTCTCTTTTTTTCTTTATTAGT
CTTGCTAGCGGTCTATCAATTTTGTGATCTTTTCAAAAACCAGCTCCTGGATTCAATTTTTTGAAGGGTTTTTTG
TGTCTCTATTTCTTTCAGTTCTCCTCTGATTTTAGTTATTTCTTGCCTTCTGCTAGCTTTTGAATGTGTTTGTCTCTGC
TTTTCTAGTTCTTTTAATTGTGATGTTAGGGTGTCAATTTTGGATCTTTCTGCTTTCTCTTGTGGGCATTTAGTGCTA
TAAATTTCCCTCTACACACTGCTTTGAATGTGTCCAGAGATTCTGGTATGTTGTGTCTTTGTTCTCGTTGGTTTCAA
GAATATCTTTATTTCTGCCTTCATTTCTGTTATGTACCCAGTAGTCATTCAGGAGCAGGTTGTTTCAGTTTCCATGTAGTT
GAGTGGTTTTTGAATGAGTTTCTTAATCCTGAGTTCCAGTTTGATTGCACTGTGGTCTGAGAGACAGTTTGTATATAATTT
CTGTTCTTTTACATTTGCTGAGGAGAGCTTTACTTCCAACATATGTGGTCAATTTTGGAAATAGGTGTGGTGTGGTGTGCTGA
AAAAAATGTATATTCTGTTGATTTGGGGTGGAGAGTTCTGTATATGTCTATTAGGTCTGCTTGGTGCAAAGGTGAGTTC
AATTCCTGGGTATCCTTGTAACTTTCTGTCTCGTTGATCTGTCTAATGTTGACAGTGGGGTGTAAAGTCTCCCATTA
TTATTGTGTGGGAGTCTAAGTCTCTTTGTAGGTCACTCAGGACTTGCTTTATGAATCTGGGTGCTCCTGTATTGGGTGC
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ATCTTTGTGTTGTTTAAAGTCTGTTTTATCAGAGACTAGGATTGCAACCCCTGCCTTTTTTTTGGCTTTCCATTTGCTTGGT
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TGGGTCTTGACCCTTTATCCAATTTGCCAGTCTGTGTCTTTTAATTGGAGCATTTACTCTATTTACATTTAAAGTTAAT
ATTGTTATGTGTGAAGTTGATCCTGTGATTATGATGTCAGCTGGTTATTTTGTCTCATTAGTTGATGCGGTCTCTTCTTA
GCATCGATGGTCTTTACAATTTGGCATGTTTTTGCAGTGGTTGGTACTGGTTGTTCTTTCCATATTTAGTGCTTCTT
CTGGAGCTCTTTTAGGCCTGGTGGTGACAAAATCTCTCAGCATTTGCTTGTCTGTAAAGGATTTTATTTCTCCTTCACT
TATGAAGCTTAGTTTGGCTGGATATGAAATTTCTGGGTGAAAATTTCTTTTCTTTAAGAATGTTGAATATTGGCCCCAT
TATCTTCTGGCTTGTAGAGTTTCTGCCAAGAGATCTGCTGTTAGTCTGATGGGCTTCCCTTTGTAGGTAACCCAACCTT
TCTCTCTGGCTGCCCTTAACATTTTTTCTTTCATTTCAACTTTGGTGAATCTGTCAATTATGCGTCTTGGAGGTGTGCT
TCTTGAGGAGTATCTTTGTGGCGTTCTCTGTGTTTCTGTATCTGAATGTTGGCCTGCCTTGCTAGATTGGGGAAGTTC
TCTTGATAATATCCTGCAGAGTGTTTTCCAACCTTGGTTCCATTTCTCCCATCACTTTGAGGTACACCAATCCGACATA
GATTTGGTCTTTTTCACATAGTCCCATATTTCTTGAGGCTTTGTTCTGTTTCTTTTATTCTTTTTTCTCTAAACTTCCC
TTCTCGCTTCATTTCAATTCATTTCACTTCCATCACTGATACCCTTTCTTCCAGTTGATCGCATTTGGCTCCTGAGGCTT
CTGCATTTCTTACGTAGTTCTCAAGCCTTGGCTTTTCACTCCATCAGCTCCTTTAAGGACTTCTCTGTATTGGTTATTC

Fig. 9.109

ATTCCTCCTTCACCCTGTTCCCATCTATGAGAAATTATATTTATTAGTGTGTCATGTTTATTTTTTTCAGTGTTTCTATAT
GTAATACTCATTATATGTTATACAAACACAAATATACATATACATAATTTCCCTTCCATTTTTATCAATTTCCCTTCCCTTT
CTCACTGTCCTGCTCCTAGCTTTAATTTTACCAATTARTAAAACATACTAAAGATTTTTTTTCATATCAGTGCATGAAGC
TTTTCTCTTTTTTTCTTTCAACTGCATAATATTCCTACTATGTGGAAATATCATGGTTTATTTAAATAAATTTCTTATCAG
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CATCATTTCTCATTTTCATATGTGTACAAAGTATCTGTGGGATAGAGTTCCAGAACTGAAATTGCTGCTTGAAAGGGTAA
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TATAAAGACACATGCATATGTCTGTTTACTGCGGCACTTTTCACAATAGCAAAGACTTGGAACCAACCCAAAAGTCCAT
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Fig. 9.112

Fig. 9.113

ATAGGCCAATAGCAGTCTCTAAAATGGAGGCAGTAATTAATGGCCTACCAACCAAAAAATCCTAGGACCAGACTGATT
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GAAATTGAGCCAGTCAACTGAACCCACCTTATACCACCATGAAACACCCAAGGTGATGAAATAGAACAAAAGAAAAAA
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GGCCGAGGTGGGTGGATCACGAGGTCAAGATATCAAGACCACCTGGCCAACATGGTGAAACCCCATCTCTACTAAAAA
TACAGAAAACCTAGCCGGGCATGGCAGTGGGCGCCTGTAGTCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGCGTGA

Fig. 9.115

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ACACGGGAGACGGAGCTTGCAGTGAGCCGAGACTGCACCCTGCACTCCAGACTGGGCAACAGAGCGAGACTCCATCTC
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GCAAAATGAATCCAGGAGCACATCGAAAACTAATCCACCACAATAAATTGGCTTTACCCATGGGAGGCAAGGTTGGTT
CAACATACACAAATCAATAAATGTGATTCATCACATAAACAGGACCAATGCAAAACCCACATGATTATTGTAATAGATG
CACAAAGGCTTTTGATAAAATTCATCACCTCTTCATGTTAAAAACCTCAACCACCTAGGTACTGAAGGAACGTACCTC
AGAAAAATAAGAGTTATCTATGACAAATCCACAGCCAACATCATACTGAATGGGCAAAAGCTTGAAGCATTCCACTTGA
AAACCAGCACAAAGACAAAGACGCCCTTTCTCACCACTCCTATTCAACATAGTATTTTAAGTCCTGGCCAGAGCTATCAG
GCAAGAGAAAGAAATAAACAGCAAGGCAATACCAACAGCCAAGGCAATACTAAGCAAGGAGAACAAAAAAGAACAAAT
CTGGAGGCATCACACTACCTGATTTCAAACCTATTCTACAGGGATACGGTATCCAAAACAGCATGATTCTGGTATAAAAA
CAGAAACATGACCAGTGAAACAGAATAGAGAGCCCTGAAATAAGGCTGCACACCTACAACCATCTGATCTTAGACAAAG
CAAAGGGGAAAGGACTCCCTATTCAATAAATGATGCTGAGAAAACCTGGCTAGCGATATGCAGGAGATTGAACTGGACC
CATTCAGTACACTATACATAAAAAATTAACCTCAAGATGGGTTAGAGACTTAAACGTAAAACTCAAAGCTATAAAAGCTCT
GGAAGACAACCTAAGCAATACCATTCAGACATAGGAAGTGGCAAAGATTTTATGATGTAGATACCTAAAGCAATTGCA
ACAAAAGCAAAAAAATGATAAATGGGATCTAATTAAATACAGAGCTTCCTCACAGCAAAAGAACTACCAACAGAGAAA
ATAGACAACCTACAGAATGGGAGAAAATATTTGCAAACTATGCATCTGACAAAGTTCTTATATCCAGCATCTATAAGGA
ACTTAAATTTATAAGAATTAAACAACCCCATTAATAAAGTGGGCAAAGGACGTGAACAGACACTTTGCAAAAAAAGTACG
TGCAGCTAACAAAGCATATGAAAAAACTCAGTATCACTGATCATTAGAGCAATGCAAGTCAAACCCACGAGATGCCATC
TCACATCAGTCAGAATGGCTATTATTAAAAAGTCAGAAAATAACAGATACTGGTGAGCTTGTGGAGAAAAGGAAATATT
TATACACTGTTGGTGGGAATGTAAATTAGTTAACTATTGTAGAAAAGTAGTGTGGCAATTCCTCAAAGAGCTAAAAACA
GAACTACCTGTCAACCCAGCAGTCCCATTACTGGATGTATACGCAAGGGAATAGAACTGTTTCATCATAAAGACACAT
GCTCACATATGTTCACTGCAGTACTATTACAAACAGCAAAGACATGGAATCAATCTAAATGCCATCAATGGTAACTG
AGTAAATAAATTTGTGGCACATATACTCCATGGCATACTATGCAGCCATAAAAAGGAATGAGATCGTGTCTTTGCGGGA
ACATGGATAGAGCTGGAGGCTGTTATCTTTAGCAAATAATGCAGGAACAGAAAATGAAATACTGCATTTTCTAACTTA
TAAGTGGGAGCTAAATGATGAGAAGACATGGACACAAAGATGGAACTAGGCTTAGCAACTGGGTGACAAAATAATCTG
TACAACAACTCCCATGACACAAGCTTACTTATATAACAACTTGCACATGTATCTCTGAACCTAAATAAATGTCTTT
TTAATATCTGGATCCAAATTGTTTTTACTGGCCAATGGAAAAAGATTTTTTGCCAGATGGCTAAATCTTTTGAATAATA
TTTGTGAAAAAGACTTTTAAAGATTTTGAATAATATTTGTGAAAAAGACTTTTAAATAATATTAGTGAAAAAGACTTCCT

Fig. 9.116

TACAGAAGGCAAATTAAGTCTTAATTTAATTTTGGCAGCTTTTAATGTGGCAATCTTTGATTCTTTTATTCTTTTAGATGG
CTGTGTGCACCAATTAAGAATGCATCCCATTTGCTAAAGAAATTAATAATTTAAAAGAAAAAATCTTAAACAATTTAA
TACTACATCTCAAGGAAGTAGAACAATAACAACAGACTAAGGCCAAATTAGGCAGAAAGAGGGAATTAACAAAGATTTG
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AAATGAAAGAAGAGAAATCACAAATGATGCCACAAAATAAGAAAGAAATCAAAGAGATAACTATGAACAATTATGTGCC
AACAACTGGATAAGCTAGAAGAAATGAATAAATTCCTAGAAACATACAATCAACCGAGACTGGATTCTGAGGAAGCAG
AAAGTCCGAACAGACCTATACTAGTATAGATATTGAAATAGTTATTACAAAGCTCCCAGGAAGGAAAAGCCCAGTACC
AGGTGGCTTCACTGGTGAAATTCTAACAAACGTTTAAAGAAAAATTTATACCATTTCTTCTAAAACTTTTCAAAAAAAT
TGAAGGGGAGAGGGAACATGTCCAAATTCATTACTCTGATACTAAAGTCAGACAAAAGCACCACAAGAAAACAAAATA
TAGGCCAATATCCCTGAAGAATGAACATACAAATTCCTCAACTAAACCCCAGCAAACTGAATCCAAGAGCACATTAAGG
GGATTATACACCATGACCAAATGGGATTTGTACTTTGGGATGTTAGAATGGTTTCAGTGTGTAAAAATTAACGTGAAATTC
CACATTAACAAAATAAAGCATAAAAAACATGTGATCATCTCAAGATACAGAAAAAGGGTTTGACAAAATTTAACATCTTT
TAATTA AAAACTCAAACCTAGTAATAGATTATGTCAAATACAAATACAAATATTATATACCTCAACATAATAAAGCCAT
ATATTAAATGCCCATCACTAACGTCATACTCAATGGTGAAAAACCGAATGAAGGCTTTTCTCTAAGATTAGAAACAAG
ACAAGGATGCCCACTTTCACTGCTTCTATTTCAGCAGGATACTTGAAGCCCTATCTAGAGCAATTGGGTAAGAAAAAAG
TAAAAAGGCATTTAAATTGGGAAGGAAGAAATAAAATTTTCCCTGTTTGCAGGTAACATGATCCTATATATAGAGAAAA
CTCTGAAGATTCAATTAAGAACTGTTAGAACTAACAAATGAATTTAGTGAAATTACAGAGTACAAAATCAACATACCA
GAATCAGTTGCATTTCTATACACTAACAACTATCTGAAAGGAAATTAAGAAAACAATGCCCATCTAACTAGTGC
CAAAAAGAATAAAATAATTTAGTAATACACCTAAACCTAACTAACAAAGGTGAAAGATTTGTGTATTAAAACTACAAAA
CATTGATGAAAGATACTAAGCAAATCAAAGACATTTGAATGTTTCATGAAGTGAAGACTTAACATTTGTTAAATATCCA
TACTACCAAAGAAATCTACAGATTCAATGCAATCCCTATCCAAATCTCAATGTCAATTTTTTTACAGAAAGTGAAAAAA
GCTCTAAAATTTATATGGAGCCATGAGTCAAGAAATAACCAAATCAATCTTGAAAAAAGAATAAAGCTAGAGACATCA
CATTTCTCTGATCTCTAGATATGTTACAAAACCACAGAGATCAAAACAATATGATACTGGCATTAAACAGACATATAGA
CCAATGGAACATAAGAGAATCCAGAAATAAATCTATGCATACTTGGTTGTTTTATTCTATTTCGGGCTACTATAATAAAA
TACCATAAACTGGGTAGCTTATAACAACAGAAATATATTTCTCACAGTTCCAGAAGTTTGGAAATTCCAAATTGAGGCA
CAGGTACTGATGGACTGCTTCTCATAGTTAGTGCCTTCTCACTGTGTCTCACATGGTGGAAAGGTGAAGCATTCTCT
CAGATGTCTTTTTATTTCATGAAGTCTCCACTTTTTATGACCTAATCATCTCTGTAGTCCCGTCTCCTAATGCCATTCAA
CATATAGATTTAGGGAGACAAAGATTCAACATATGGATTTTGGGTGACACCAACATTTCAGTCTATAGCAACAGTCAACT
GACGTTTGATAAAGGTGTCAAGAATACACATAGAGATTGGAGAGTCTCTTCAACAAATGGTACTTGGAAGACTGGATAT
CCATATTAAGGAATGAACTGGATGCTTGCTTACATCATATGCAAAAATCAACTTGAAGTGGCTTAAAGATTAAACAT
AGACCAAGACTATAAACTACTAGAAGAAAACCTTCATGACATTGGTCTTGGCAATGGTTTCATGGATATGACATCAAA
AGCACAGGCAACAAAAACAAAAGTAAACAAATAAGAATACATCAAACCTAAAAAGCTTGTGGGTGAATTGTTTGAGCTCA
GGAGTTCGAGGCCAGCCTGGGCAACATAGTGAAACCTTGTCTCTACAAAAAATTAAAGAAAAAATTAGCTGGATATGGT
AGGCTCACACCTGTAGTACCAGCCACTTGGGAGGCTGAGGTAGGAGGATTGCTTGCACCTGGGAGGTGCCAGAGGTTGC
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ATTAATAAAATAATGATATGAATCAACAGAGTGAAAAGGACACTTACAGAATGGGGGGAATATTTGCAAACCGTATATC
TGATAAAGGGTTAGTATCCAAAATATCTAAAATAAACCCCTTCAGGTTATTAGTTAAAATAAAAGAGAGAGAAAAATAA
ATAAATAAATAAATAACCTGATTTAAACATGGGCTATGGACTTGGATAGACATTTCTTCAAAGAAGACATACAAATGGC
CAACAGATATTTTAAAGAAATACTCAGTGTCACTAATCATCCAGGAAATGTGAATTAAACTATAATGAAATATCACTTA
ACACCTACTATAGTAGAATGGCTACTGTTAAAAAAAACAGAAAATAGCAAGTGTGACGAGGATGTACAGAAATTGGAA
CCCTTGCACACTGTTGGTGGAAATGCAATATGGTGCAGCTTTTGTGGGAAACAAATGAAGTTCCTCTAAACATTTAAAA
ATGCAATTACATGATCCAGCAATCCCCTTTGGGGTGTTTTATTCAAAGAATTGAAATCAGGATCTCAAAGATGTATTA
GCACTCCTATATTCAATTGTAGGATTATTCACAATAGTTAAGATTTAGAAACAACCTTAAGTGTCTATTGACAGATGAATA
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TCTAGTTTATATTTGCAAAGGTGTGTCTGGGCCAGTCCAATCTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT
ATTACAGATCACATGCCGAACCTTTAGGGGAGAGGACAAATTAAGGATTAACAATGAAAGATATAGAACTGCTCAGGA
TATTTTCTAGGTTTAGGAGACTTGAGGAAAATGAATAAGTGAAATGTAGACATATATGAATGAATATATCAAACATCAG
AGGAGAAAATTTAGCTGTGTCCAGAGAGGACAGAGATGCTATCTTTCTATTGCTTGTTTTTGAACTTTCTTTCTGTTATTT
TATTATTGTATTGTTAATAATCATTGAAGTAGTGTGAGAAATTTAGAGAAAGTAGAAGTAGAGCAGTTTCTCTGAAAAA
CTTATAGATGGTGACAGATGACCACATAATTGTGTGACAGTAAAATGCTGTATTGAGCTTCTGCTGTATTATAGAAAG
AACATATGAATAGATAAAAGCACAAACAGTTTTTCTTTTCACTATGTCTCAGGAGAAAACAATGCACATCGGCATATATG
GAAACAAATGAGATTAGTTAAAAGAAGGAATTTAAATTATGTAGACTAAGCCAGGGTAATGCTTACGAATGAGATGACT
CATCTAATAAAAGCATCAAAGAGAAAAACTGTTGTTTTAGTTTCAGATTTTCTTGGTTCGTTTTCTCTGAGATGATGATGAT

Fig. 9.117

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GATGATGATGATGATGATGATGATGATGATAATTTGCTTTAGGAAGGAGCTGCACACTACAAAATGCAACAGAGTTGAT
AATGATAAGATTAGTTGAATTAAGTAATTCTAGACCCCAAATTGATTATTTTCATTTCACTTTCTAGGTCAATAGTGTAC
CTTGTTGCAATGAACTGAGTTGGTTGTAAAAAGTCTACTATCTTATCAAAAGATTGTTCAAAAACATCTCCATGTACTA
AATACTATCAAGCAAAAATATATTGTGTATAGGGAAATACAGTATTTAAAAAAATTGGTAGAATCTACAATGAACTCAA
ACAAATTTACAAGAAAAAAACAAACAACCCCATCAAAAAGTGGGCGAAGGATATGAACAGACACTTCTCAAAAGAAGAC
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AAGACACACGCACACGATGTTTATAGCTGCACTATTCACAATAGCAAAGACTTGAACCAATGTAAATGTCCAACAAC
GATAGACTGGATTAAGAAAATGTGGCACATATACCCATGGAATACTACGCAGCCATAAAACATGATGAGTTTCATGTCC
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CACTCATAGGTGGGAATTGAACAATGAGAACACATGGACACAGGAAGGGGAACATCACACTCTGGGTACTGTTGTGGGG
TCGGGGGAGCGGGGAGGGATAGCATTAGGAGATATACCTAATGCTAAATGATGAGTTAATGGGTACAGCATACCAGCAT
GGCACATGTATACATATGTAACCTGCACATTGTACACATGTACCCTAAAACCTTAGAGTATAATAATAATAATGAT
AATAAATTGGTAACCTAGTTTAAGGTCATGTAACCTACTCACACAAATAATCACATAAGATGTAATATAGAATGTGCAT
GTAAACATAAATTTTATACATAAAATAAATAATAGATCATAGCCCTGATGCATGCGTGCACAAACACACACACACA
CACACACACGCACACACACACTTGCACCTATATTTGCAATAGCCTAGCTACTAAGGAATCTGATTAAACATAGATTCTATC
TTGCATATCCGATTTCTTAATAGCAGAATTAAGAGCTTCTTGGAAAGAAAAGGCTTATACTCAAAATCTGCTTTTCTGAA
TCCTTTTGAGTTGAGCACAAAAGGAAACCAAAAAGACCCAAGAATAGCAATGCAATGTGTGTATTGAAGGTTTGTAGG
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GCCCTTGAAAAATCAAGCAAATAGATGGCTTCACATTTCAATTATCTAACCCCGATATGGCCTGACATTTTATACTGGGAG
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TGTGTGGATTTTTTTCTAATTGAAAACAGCACAACGAGAGCTCTGAGTATGAGATGTAGTTATACAAAAGGCATAAGA
TGCTAGACACCTTCAAATACTGGGTGTTCTCCTTACTAGCCATATGACCTTAAGCAATGTATTTAACCTATTCTTTGTT
CTCATTTTCTCTCATTCATAAAATATGCACAATAAACTTACCTTATAGTTTTGTTTCAAATATAAAATGAATAAATATT
GTTAAATTATTTTCGAGCATTCTTTGGTACACAGTAAACCCTCCGTAAGTATTAACCTTTTATGGATAAAAACCTTTTAAA
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ACCTCCATGCAAATCTCACTTACCTACCTCTTTATCCATCAAAAATAATCTATTAATAATCAGTTATACTTACTTATTTA
GCACTTAAATATTTTAGACTTTGTGCTAAGTGCCTCACATATAGCATTTTTTACTCTTCCAGATAGCTTTCAAGGGAGCT
TTCATAACAACCATTTTAGCATTTGTTTCGATCTCATTCTCAATCTCCATCTCTTTCAAACATAAACTAAATTTCTCCC
TTGGCAAGGTTGGCCTACACCCAGGAACAAGCAAGGAAAGCCACCCTCTGAGGCTAGAAGCAAGATGGAGTCAGCCATG
CTAGCCTTCTCTCATTGTTATAATCTTTGCAAAGCTGGTTTTCATATTTTACTAATCTTTCTCTGTAAATGGACAAACA
TCCTTCTTGATGCTACTTCACACACCTTTTCCAACCTCTCTGTACTTGACACAATTAGACAAAGCATTTGCTCAATAAA
TATTTATAAAGTTAAGAAAGAATCACTTTATACCTAAGAGTGGATCCTAATGTTTACATAAATATATTCCTGTATTGAG
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CTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTTCTTT
GATACATATACAGAACATGCAGGTTTCTTACACAGTATATGTGTGTCACGATGATTTGCTGCACCTATCAACCCATCAC
CTGGGTTTTAAGCCCTGCATGCATTAGCTATTTGTCTTGATGCTCTCCCTCCCTCGTGCCCCCATGACCCAACAGGCC
CTGGTGTGTGTTGTTCCCTCCCTGTGTCCATGTGTTCCCATTTGTTCAACTCCCCTTATGAGTAAGATGGATTTCTGT
TTGCTTTTCTGATCCTGTGTTAGTTTGCTGTGAATGATGGCTTCCAGCTTCATCCATGTCCCTGCAAAGGACATGAACT
CATTCCTATTTGTGGCTGTATAGTATTCTATGGTGTATATGTACCACATTTTCTTTATCCAGTCTATCATTTGATGGGCA
TTTGGGTTGGTTCTATGTCTTTGCTATTTGTAATAATGCTGCAATAAACATACATATGCACGTGTCTTTATAATAGAAT
TATTTATATTCCTCTAGGTATTTACCCAGTAATGGGATTGCTGGGTCAAATGGTATTTCTGGTTCTAGATCCTTGAGGA
GTTGCCACACTGTCTTCCACAATGGTTGAACCTAATTTACATTTCCACCAACAATGAAAAAGTGTTCTCTTTCTCCACA
GCCTCTCTAGCATCTGTTGTTTCATGACTTCTTTTTTAATAAGCGCCATTCTGACTGGTGTGAGATGGTATCTCATTGTG
GTTTTGATTTGCATTTCTTTAATGATCAGTGATGTTGTGCTTTTTTTTTCATATGTTTGTGGCCACATGAATGTCTTCT
TTTGAGAAGTGTCTGTTTCATGTTTTTGCCCACTTTTTAATGGGTTTTCTTGTGTTTTTCTTGTAAATATGGTTAAGTTC
CTTGTAATAATTATGGATGGTAGCCCTTTGTCAGATGGGTAGATTGCAAAAATGTTCTCCCATTTCTGTAGATTGCCGTTC
ACTCTGATGACAGTTTCTTTTGCTGTGCAGAAGCTCTTTAATTAGATCCCATTTGTCAATTTTAGCTTTTGTGTGATT
GCTTTTGGTGATTCCATCATAAAATCTTTGCCCATGCCTATGTCGTGAATGGGATTGCCTAGGTTTTCTTCCAGGGTTT
TTATGGTTTTTGGGTTTTACATTTAAGTCTTTAATCCATCTTGAGTTGATTTTTGTATAAGGTGTAAGGAAGGGGTCCAC
TTTCAGTCTTCTGCATATAGCTAGCCAGTTTTCCAGCATCATTTATTAAACAGGGAATCCTTTCCCATTTGCTTGT

Fig. 9.118

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TTGTCAAGTTTGTGGAAGATCAGATGGTCTTATATCTGAGGTCTCTATTCTGTTCCACTGGTATATGTGTCTGTTTTGA
TACCTGTACCATGTTTTGGTTACTGTAGCCTTGCAGTATAGTTTGAAGTCAGGTAGTGTGATGCCTCCAGCTTTATTCT
TTTTGCTTAGAATTGTTTTGGCTATACGGGCTCTTTTATGGCTCCATATGAATTTTAGAGTAGTTTTTTTCTAATTCTG
TGAAGAAAGTCAATGGTAGTTTGATGAGAATAGCACTTAATCTATAAACTACTTTGGGCTGTATGGCCATTTTCATGAT
ATTGATTCTTCCTATCCATGAGCATGGGATGTTTTTCCATTTGTTTGTGTCTCTCTTATTTCTTGAGCAGTGGTTTG
TAGTTCTCCTTGAAGAGGTCTTCATGTCCCTTGTTAGCTGTATTCTAGGTGTTTTATTCTCTTTGTAACAATTGTGA
ATGGGAGTTCATTCATGATTTGGCTCTCTGCTTGTCTATTGTTGGTGTAAAGAAATGCTTGTGATTTTCACACACTGAT
TTTGTATCCTGAGACTTTGCTGAAGTTGCTTATCAGCTTAAGGAGATTTGGGGCTGAGACAATGGGGTTTTCTAAATAT
AGGATCAAGTTGTCTGCAAACAGAGACAATTTGACTTTCTCTTTTCCCTATTTGAATACGCTTTATTTCTTTCTCTTGCC
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CAAAGGGAATGCTTCTCGCTTTTGGCCATTGAGTATGATATTGGCTGTGGGTTTTGTCAGAAGTAGCTCTTATTACTTTG
GGAAATGTTCCATCAATATCTAGTTTATTGAGAGTTTTATCATGAAAGGATGTTGAATTTTGTGCAAGGCCTTTTCTGC
ATCTATCGAGATACCAATTACCAAAATTCTAAGATTACATGGTGTCTGCACTGAATAGATCTATGCAAACCTACCCCCA
AAGTTCAAGGAAGCTGAGAGGCTGAAGAGGCTGACATATCCAGCTTCTCAGGAAAAATGAAAAACAAAAACATTTAA
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AGTGAAGGAACATGTAGGACAGTTGAAATCAACTCCTCAGGGAAAGGCAAGCATGCTATGTAAATCTGCCTAAGGGAAG
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CTGGAACCTGGGGTAAATCAGATGTCAGTATGGTGGACTGACAATGAAGATGGAGTTGCTTTAGTCTCCACACATGGGAA
CAGATAGAAAAAGGTGTCTTTATAACCAAAATGTTGGCTTTTACAAAGCTCTGCTGTAATTTCTAAATAGATGGAAGAGG
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TAAATCTCTCATAGAATAATTATTAGAGTATTTATGAATTTTCTTCTCTTCTCCAAATTATCTTCACTTGGGGTTATTT
TATCTTTATTTGTGTATCTTGGACTGTCTTTTTTCAAAGCAAGCTTTCTTTAGATGCCTGATGAGCTATGATTATAATT
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GCAGGGTTGGATAGTGGCTATTTCTACCCATAGATACTTGTAAATTGTCAGTTCCCCATGTCACCTCCTGTACCCTTTCT
GTCTTCATTGCATGTGGTGTTTTTGGAATTTCTTAGCCTTTCTGGAAAGGCAATTTCTCTCTCTCAGTCTGTCTATTTAT
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TGAGTGACAATACTGCCTTTTAAATTTTTTCTCAAATGTTATAGGGCAACAAGAGATTGGCAGTATGAGCTGTCTTTTT
ATTGTATTTTAAATGGTAACTTGTACATGATTATGGGGATATTCTAACTTTTATGATTTTTTAAACAACCTCTCAGTTCC
ATCCCATAAAAGAGCTTTATGTAGAAATACTTTTAGCAGCTTTTAAAGTTTATTTTCATCTTCTCTTCCCAGGAGACAAG
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ACAGTGCTAACACTGAAAAGGAAAAGAGTAAGATTTTGTAAATTTATAAATAATCCTTATGGGGATGTTATTATTAACT
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TATTTCACTGTTTTTAAATCATACTCAGGAAGGGTTTTAGTGAAGAAACAGGAGTGAGATTAATCAAATAGCTCAAAT
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GTACCTTGCTGTTCTTGGGTTTGAAGGATCAGTATTTTCTTGCCATTTATCACACTTAAACATAGAGCATCCAGA
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TGTGGCTGGAGCTGAGTATTGAAATAGGAGAAATATTTGTTTCAGATATTACTCTTTTTTATTTTTTCATGACTGAATTCC
AGGCATACGTACCTTCAAATACCTTATTGCAGATGACCTAGAAGGCTGCATGTAAAGATAAATAACATCCTAACATTTGT
ATATTGTTGCTTCTCAAATTTCAACACTGTGATTACTTGCTGTGTTGGTTACTGATCTTTATTTTTCATTTACTTAGCAT
CTTCATTGATAATCCTGATTGATGTAGTTTTACTTTTCATCATCTAAATATCTCTAGTTTCTAATTTATTCTGTGTAGAG
ACTTTAGTTTCATCAAATATAAGTGGGCATGAAATTATATACCTGAGCCAATAGAAGAACTCAAGACTTTTATTTGCC
TAATGTAGGATTTAAAAACCAATTCAAAAACAATTGTTTATATATGTGGGAAAATTCCTTTAAGTTTCTGTTGATAGGT

Fig. 9.119

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ATTGGAATAGGTTTCCTAGTAAGATATTAAATGTGTCATCCAAATAAATTTTTAGGTTACATAAACACTGATCTTGAAT
AAAATAGCTATAGTTCTACCTGGAGGTAGAGAGTCATAATAATATCAAATTAGAGCTCATGTTTGTGTGTGTATGTGTT
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GCAGAGTCTCAGTGAAAACCTGTTTGGGGGTGGTGCCTCTGATTCTCAGTGCAGCCTGCCAGCCTCATTGGGAAAAGG
TGTTTAATATAGGCAGCGATTTTCTGTATTATGAACTGAAATAGAAAATGCTGCGATGGGTTCATTCAAGGGAGACCCA
ATATCTTACTTACAATGAAGAAATCAGGCAATTTTGGGGAGTAGAAGTGGAAGTGTGACTCATACATATATTGGTATTT
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GTAGTGGTTAAGAGCATAGGCTCTAAAGTCAGACTGCCTGGCTTTGAATCCTCTCTACCACTTACTGACTGAGTGACCT
TGGATTGAGTCACTGAACTTCACTTTTCAATTTTTTATGTCTTATTGGATAAGTTTCAATCAGAGTTGAGGATGAAGTG
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CATTTATGGTTATTCCAATATATACATTTTGTGTAGAAGGCTTAGTCTTTCAATTACAATAAGCTATTGGTATAGTTTA
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ACTTATTTTTTTTTTATTATACTTTAAATTTCTGGGATACATGTGTAGAATGTGCAGGTTTGCTACATAGGTATAAACATA
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CCCACCGACAGGCCCTGGTGTGTGATGTTCCCTCCTGTGTCCATGTGTTCTCATTGTTCAAATCCCCTTATGAGTG
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GATGTGCTGCTGGATTCTGTTTGCCAGTATTTTATTGAAGATTTTTGTCATAGATATTCATCAGGGATATCGGCCTGAAA
TTTTTTCGTTGTGTCTCTGCCAGGCTTTGGCTTCAGGATGATACTGGCTTCATAAAATGAGTTAGGGAGGACTCCCTCT

Fig. 9.120

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TTTTCTATTGATTGGAGTAGTTTCAGAAGGAATGGTACCAGCTCCTCTTTGTACCTCTGGTAGAATTCAGTTGTGAATC
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GCTGTGTCTAGAGATTCTGGTACATTGTATCTTTGTTCTCACTGGTTTCAAAGAACTTATTTATTTCTACCTTAATTT
CATTATTTACCCAGTAGTCATTCAGGAGCAGGTTATTCAGTTTCCATGTAGTTGTGTGGTTTTGAGTGAGTTTCTTAAT
CCTGAGTTCTAATTTGCTCTGTGGTCTGAGAGACTGTTTGTTATGATTTCTGTTCTTTTGCATTTGCTGAGGAGTGTTT
TACTTCCAATTATGTGGTCAATTTTAGAATCAGTGTGACAAGGTGCTAAGAAGAAATGTATATTCTGTTGATTTTGGGTG
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CACACCACACCTATTCCAAAATTGACCACATACTTGGAAGTAAAGCTCTCCTCAGCAAATGTAAAACAGAAATTATAAC
AACTATCTCTCAGACCACAGTGCAATCAAACCTAGAACTCAGGATTAAGAATCTCATTCAAACCGCTCAACTACATGG
AACTGAACAACCTGCTCCTGAATGACTACTGGGTACATAACGAAATGAAGGCAGAAATAAAGATGTTCTTTGAAACCA
ATGAGAACAAGACACAGCATAACGAAATCTCTGGGACGCATTCAAAGCAGTGTGTAGAGGGAAATTTATAGCACTAAA
TGCCCAACAAGAGAAAGCAGGAAAGATCTAAAATGGACACCCTAACATCACAATTAAAAGAACTAGAAAAGCAAGAGCAA
ACACATTCAAAGCTAGCAGAAGGCAAGAAATACTAAAATCAGAGCAGAACTGAAGGAAATAGTGACACAAAAAACCC

Fig. 9.121

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TTCAAAAAATTAATGAATCCAGGAGCTGGTTTTTTTGAAAGGATCAACAAAATTGATAAACCGCTAGCAAGACTAATAAA
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ATCCCTGGGATGCAAGGCTGGTTCAATATACGCAAATCAATAAATGTAATCCAGCATATAAACAGAACCAAGACAAAA
ACCACATGATTATCTCAATAGATGCAGAAAAGGCCTTTGACAAAATTCAACAATGCTTCATGCTAAAACTCTCAATAA
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CAAAAATGGAAGCATTCCCTTTGAAAATGAGCACAAGACAGGGATGCCTTCTCTCACCCCTCCTATTCAACATAGTGT
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GGAGCCATTTGTTGTTACCCACATTTTGAAGTCTACTTCTGTCAATTTGTCAAATTCATTCTCCATCCAGTTTTGTTC
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TTTGTGACGTTGATGCTACTCCTTTCTGTTTGTAGTTTTCTTCTAACAGTCAGGCCTCTCTGCTGCAGGTCTGCTG
GAGTTTGTCTGGATGTCCACTCCAGACCCTTTTGCCTGGGTATCACCAGCAGAGGCTGTAGAACAGCAAAGATTGCTGC
CTGTTCTTTCTCTGGAAGGTTTTTCCAGAGGGGCACCCACCAGATGCCAGCTGGATCTCTCCTGTATGAGGTGTCTG

Fig. 9.122

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TCGACCCCTGCTGGGAAGTATCTCCAGTCAGGAGGCACGGGGGTGAGGGACCCATTTGAGGAAGCATTGTGTCCCTTA
GCAGAGCTCAAGCACTGCGCTGGGAGATCCACTGCTCTCTTCAGAGCCAGCAGGCAGGAATGTTTGTCTGCTGAAGCAG
CGCCTACAGGCACCTCTTCCCCCAGGTGCTCTGTCCCAGGGAGATGGGAATTTTATCTATAAGCCCCTGACTGGGGCTG
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GCTTTTTCTGTGTGATGAGCTGTAAACCTTCATATTAGAAAACTCAGAAAAGAATTTTGCTTAGACGCTAATCAAATA
CAAAAATTGTGGCTGATGGAACTACACATAGATAAATTAGTCCAATATTCTTCTACTTGTGAAATTTAAATAACTTCA
TCTTAAGAAATAAAGGTAATTTGGGAAAATTTGAAAAGGAAGTGTTCAGTTTTGATAGAGTGAATGGGGCATTCAAAT
ATATTTCTAAATGAATGATAAAAATAAAATACTTGCAATATTATCAGAATTTATTTCCACAATAAATATGTTTCAGGAAT
GTTAAGTGGGGAAGCCTGGGCAGCCATTGTGCTGTAAGTGGCTCTGTTTATTTTTTTGCTCTTCAAAGAGAGAAAGAAAGG
AACCTTAGCACAATAGCAGAATTTCCCTGATGGTGGCTATTACAATTTTACCACTGAGGAAAGGACATCTATGGGTCTTT
GAAAAGCTAGGAAACATCTTGAATATCAGAAATTGTAAATGAATACTACTTGGTGTAGATTAAACTAAAGACATGGGAT
TATATGGTCTCAAAGCCACAGATGTAGTTGGGTCTAAGAACCCTTGTATATATCTATATTTTTTTTATTTTTTTTTTATT
TTTTTTCTTTTATTATTATTATAGTTTAAAGTTTTAGGGTACATGTGCACAATGTGCAGGTTAGTTACATGTGTATACAT
GTGCCATGCTGGTGTGCTGCACCCATTAACTTGTCAATTTAGCATTAGGTATATCTCCTAAAGCTATCCCTCCCCCTCC
CACCACCCCAACAGTCCCCAGTGTGTGATGTTCCCTTCTGTGTCCATGTCTCATTTGTTCAATTCCCACCTAT
GAGTGAGAATATGTGGTGTGTTGGTTTTTTGTTCTTGTGATAGTTTACTGAGAATGATGATTTCCAATTTTCATCCATGTC
CCTACAAAGGACATGAATCATTTTTTTTTATGGCTGCATAGTATTGCATGGTGTATATGTGCCACATTTTCTTAATCCAG
TCTATCATTCTTGACATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATAGTGCCACAATAAAACATACATGTGCAT
GTGTCTTTATAGCAGCATGATTTATAGTCCCTTTGGGTATATACCCAGTAATGGCATGGCTGGGTCAAATGGTATTTCT
AGTTCTAGATCCCTGAGGAATCGCCACACTGACTTCCACAATGGTTGAAGTAGTTTACAGTCCCACCAACAGTGTAATA
GTGTTCTTATTTCTCCACATCCTCTCCAGCACCTGTTGTTTCTCTGACTTTTTTAATGATCACCATTCTAACTGGTGTGAG
ATGGTATCTCATTTGTGGTTTTGATTTGCATTTCTCTGATGGCCAGTGATGGTGAGCATTTTTTTCATGTGTTTTTTGGCT
GCATAAATGTCTTCTTTTGAGAAGTGTCTGTTCATGTCTTCACTCACTTTTTTGATGGGGTGTGTTGTTTTTTCTTGT
AAATTTGTTTAAAGTTCAATGTAGATTCTGGATATTAGCCCTTTGTGATGAGTAGGTTGCGACAATTTTCTCCCATTT
TGTAGGTTGCCTGTTCACTCTGATGGTAGTTTCTTTTGTGTATATGGAACATAAAAAGAGCCTGCATAAGCCAAAA
GAACAAAGCTGGAGGCATCACACTACCTGACTTCAAACATACTACAAGGCTACAGTAACCAAAACAGCATGGTACTGG
TACCAAAACAGAGATATAGATCAATGGAACAGAACAGAGCCCTCAGAAATAACGCCGCATATCTACAACATCTGATCT
TTGACGAACCTGAGAAAAAGAAGCAATGGGGAAAGGATTCCTATTTAATAAATGGTGTGCTGGGAAAACCTGGCTAGCCAT
ATGTAGAAAGCTGAAACTGGATCCCTTCCCTTACACCTTATACAAAATCAATTCAGATGGATTACAGACTTAAACATT
AGACCTAAAACCGTAAAAACCTAGAGAAAACCTAGACATTACCATTACAGGACATAGGCATGGGCAAGGACTTCATGT
CTAAAACACCAAAAGCAATGGCAACCAAGCCAAAATTGACAAATAGGATCTAATTAAACTAAAGAGCTTCTGCACAGC
AAAAGAACTACCATCAGAGTGAACAGGCAACCTACAAAATGGGAGAAAATTTTCGCGACCTACTCATCTATATTTTTT
AATGGATTAAACCAGCAAGAAAGACCATGGAGCCACCTAAATTTTTCCAAAATCTGCAAAATGAAGGTGATATAAATAT
GTCACCTAGACAATATTTTATTTTGTATATAAATTTTATTTTTTATCAGTTTGATAATATGCATAGGACTAAAAAATG
CAGTTGTCTTAAATATTTTAGGGTTGCTTAGGAAATCACTTTAAATAAATAAAGTGTGAGAATAAAGTTGTCTGCTT
ATGTTCCATATTTCAGAACTACCTTTGATTTCTTTAGCTAAACAATATTGCACATAAATAGGTTAGGAATTATAGACTT

Fig. 9.123

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TCAGATGTGCTACCTGTCTCCCTTCAAAACACTCATTAGGTAATGTATTAATTATTTTATAGAATCACCTGTCTTCTCT
GCTTCAGCAAAAACCCACCAGGGCAGAGATCACTCTCCCTCTTGTGTTTGAATTTTATGACTTAGCACAGTGAT
TGGCACATAAACATTATTAAAACCAATGAATGAGGAAAGCAACGAATAAGTAAACAAATGGGCCAGACTAAAAAGTAA
GTTATATGTATTAGTGAGAGTCAATAATAAATTAATGAAATAATCTTTTATTTGAAACTGAAGCAAATTAAGGGAAATT
TATGAACCTTAAATCTTTTTGTAAAGATCTAGTACCAATAAGATAAAAAAATGCTCATCCTTTTTTCCATTGAAATGT
TATGTTCAAATGAGCTTTGTTTCTATACTTTATACATCAATTAAGCTGAATTCATAGGTGCTAAGCATTTTAACATATT
GTATTGCATGTAATATTCCCTAATGCCCCAACTTCAAGATTATATAATGGCTTACTCTCTCCCTGTCCCTACCCACCA
GATAGTGTATCCACATACATTCTCATCTAGTTTTGTTCTGTGATGAAAAACCATATGCGTATCCTATTCTATGTGAA
TTAACCTGGCATGCAGGTAAGTGAAGTGCATGTGATCTGGTCTCTGCTGACATATCACAATGGGCCCCCTCCTTGCAT
GGAGGTAGCGTTTTTTTATAAGACAAAATGTTTTTAAATAGAACACATTTTCAGATTTTCAGATTTTATATGTATTTTGTGT
TTTTCTCCCTCTACCCCTTTCCAAAATTATGAATGAAATTCTAGGACCATTATAGACAAAGCACAGTTTAGTCCGAG
TAAACCAATGGACCCTGAAGTGTCCAAGTCAGAGTTTATTTGCAGAATTTAATTCATTCACACAGAGATAGGCAACCCA
GGCCTGTGAGGACACGAGGGTAAGTAGGCAACAAAGTGCCGTAGTCAGGCTTGTGTTTGTCTTTTTGGTAAGAGGACAAC
ATTGACTTCAGTGTGAGGGCATAAAGGAGACTCAGGACTTATTAATTTTTTCCCATTAATTCTGTGAACTTTGTGAAT
TCCCTAATATTCTTTTAAACAAGAGTTCGGAGACATGAGTTTATGTGCCTTCTTGGATATATTACAGGGAGTTTGCAGAG
AAAGTTGAATAAATTATTAGGTTAATGGCCTGTGTAAATTCACCAACACCTTTTCACTATCTCATCAATCATCCTGTTT
GAGTTATTGTGATTCACTCTGCTGATGAGCTCACACCCTTTTTTCTGATACAGGAATTATTGTACACCAGGGGACTGG
ATTTTAAACAAAACGTATTCCTTAGAATAACTTGAACAATGGATTGGTGGGTCTTACACTATTATGTGCTGTGTAGCT
GTACAAGTGTGTCTGCATGAGCTTTAGGACATTATTTGAGATATTTTAAAGCTATGTGTACCTCATGAACCTTGTAGCTGA
TTTTCTTAGTTCTTTTAAATATTTTCTCAGAAAACCAACAGTAAAATCTATCAGGTTTCACATGAATACACTCATTTG
TGTCATATCAACCCAAAATGAATATGATCTTCCAGGTAATGATGAAGGATGATAACTATAATTTCCAGCCAACTTT
ATTTTGAAACATCACTCAGTGTTCACATGTTTAGTGGCTGTTAAATCTATATGTCTAAGCAAACCTGTGAAGAGCATA
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CATCTCTAATTGCTTTGGCAATCCAATTATTGAGACAAGTAAGTGAGCCCATCGGGTGCCTCCCAAGGAATTAACCTGG
TCATTTGATATAATGTTGCACTCTCCCTATTTGGGAGGAGGGGTGGCCACCCTCACCTCTGCCATTGAAGATTAACCCA
CTACAAATTTTCAAAAATATAATATCTAATTTATTACACAGACTATCATATGGGTTCATAATCCTTAAGTATCTCTAA
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AATCTGTAAAACAGAGACAAAATGCCTTCTCAGAGGGTGAAAATAAGGATTAAGTACAATAATGAATGCAAAATACC
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GGTTCAAGCAATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCATACGTCACCATGCCCGGCTAATTTTTG
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AGCCTCCCAAGTGCTGGGATTACAGGTGTGAGCCACCGCACCTGCTGCATGCACATTTTGCCTGGATGATCTTCCAG
GCTACCATGTGGATCTCTCTCATAACAGCAGAGGGAGAGTTGTACAGCAATGAGGGCTTCAGAACTGTGTCTGACCCATG
CTGATTGCTCAGTGCCCTGAGCACAAGTTTTTAAACATTTTGAATCTCACCCATAATATTTCTCATCTTGCAGAGACA
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GTATTACTGCCTGTGGTAGTCTGAATAATGGTCTCAAAGATATCCAAGTCTTAATTTCCATAACCTGTGAATATGTA
ACCTTATATGTCAAAGGAACCTTGCAGATGTGGTTAATTTAAGGATCTTGAGATGGGGGGATTTCCTTGATGATCCA
GGCAGGCCCTAAATGTAATTATAAAGGGGCCTTTTAAAGAGAGAGGCAGGAAGGTCAAAGGCAGAAGAAGGCAATGTGAC
AGCAGAAGTAGAAATTGGTGTGAAGCCGCAAGCCAAGGCATGCAATGCTGAAAGCCTCTAGAAGCTGGAAGAAGAAAGG
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TCAGGACTTCTGACTTATTTGTGGTGTTTTAAACCACTCAGTTTGTGTTAATTTGTTGTAAGGGCAATGGGAAATGAAT
GCACTTCCCATCCATAACCTGTCTCTAGCTATGTTGAAGTCTTTACCATCCCCCAATCTGACTTTTACTTGCATGCT
TCTTGCTTTTACATATCTGCTTTTCTTCTGCTTCTTCTCTATCTCCACATCATTGCCCTTTCTCACAACCTTCTCTACT
CAGCAAATTCCTGCTTATCCTCTGAGAGGGAGGACAAGATCATCTCAGATATAGTTTGCTAGCTAAGAAAAGAGTCTTG
AGACTGCTTTGAAATCTTGTGGTCTTTTATCCATAGCTCATGTAAATAACTCTGAGCTAGCAGTGTATATTATTTTAT
TTCATCCTGGAACCTATGTCTTAGCAATATTTTTATTGGAGAAAAAAGTGAAGCAAACTCTCTGTTTCCACATGCCTA
TTGCCAAATTTCTCTGGGTGCTTTACTTTCACTGCACTGATTCTAGAAATTGTATTGACAACTGTGTTCTAAGATGACC
CTTTGATTCATCCAGATGGTGTGAAAATGTTCTTCTTCAAAAACCCACACTGCTTTTCAACATCCTAATGGGTAGAAT
CTCTAAATCCTTAGGGTGTCAACAAAAGCAAAGGGCATATTTACACTTCAGTTGGGTGAAATTAATTAGCATGGAAGTT
AATATTTACAATTAATGGAGATTACCTGACTTTAATAGTGAGATTGTGTAGCTACTTTGTATCACTCACTCTTTTTTTG
TTGTTGTTTTTAGAAATTGTGGCAAAAAATGGATAAAATAAAATTCACCTTTTAAACATTTTTTTGAATTCCTGGAGTG
TAAAAGTGGTAGGATAACCAAAACATCCTCTTTCTTGTGTAACCATCAACAATTGTCTTTTGAAGAGAGTTTGAATAT
CTTTTCTTTGAAAATCCTCGTGCAAACTTCACACCATGATCATTTATGAGGTAGTTATCAGACACTGGAGATGAATTA
CTGGGGATTTTTGTGTTCTGGTTCTCAGTATACAACCAAGGTAGAATATTGACATTGAAAATAAATGTCACTCGTTTC
TCAAGGAAAATTTGTAGTTTACCGAACAACAGCACAAATGAAAGCTGAGATACTTTACCAGTGATATTGTAGGTCTCAG
TGAGTAAAAACTCAATTAAGTATATTGGGGTGGCGGGGGCACTGTGTATAGATAGACCTGGACATGATCTCTAATCA
CAAATGCTACTTTTGAAGGGCCAGATTCTTGATACAGAGACATTTTATTTGGTGGCAATAACCATGGCTTGTCCACAG
AATGATGCCGTATTATTCTCCTGACCTAACTTCAAAGAAATAAAGAGTTTGAAGAAGAACTGCAGTTCTTCAAAGTAC

Fig. 9.124

GCAATATGGATTTCCAAGATGAATGTAGTTTCTCTCTCTGAGGAATTCTGAACAGTGGTAAAGTTTCACAAGTTTATGC
ATATCTTTTGCTCTACATCCTTCCCTAAAAGAACTTGTGGCAACAAACAAAGGGAAAGAAATAACATTTTTTAATAACT
ATGTGGAAAGCTCTATAGTAAGTGCTTCACATGTTACCTCATTTAATTCTTACAACCTACCTTATCTAGTAAATTATCTC
CCATTTGACACTTTAAGGAAGCAGCTAAGAGATGTTTCATAACTTTCCCTAGAAAAGGTAGGATTTGAATGCAGGTTTGT
ATTATTCCAAAGCTCACAAATGTGCTTTACGCAACATCAAAGTAACATATTGCGGGAATGAGTACCTTTCCCATTTAAAA
CAAATGAGTCCTGGAAACTCTTACCCTGTTTAGTTATGGAATGGCTCAGAAAATAGAAAGTGTTGAGATCATCAAAGAG
AGAAGTTAACAAAGAGCATTGTAATCCAGAAATAAGAACGCAATAGAGAAGTAGAAGTTGTGTGGCTAATTTTACCAA
CTAAATAGCCTGAATTATTAGTGACTATACACATTGATCAAATTAAATGAGCATACCATAGTCTAAAGGGGACGAG
ATTTATATTCTATCCAAGAAGTCATTAATTATGTTTGTACTATCTTCATCATGGTTATCATTTTTCTTAGACATAGCCT
AATCTATAAGATTTTACTGTATTTCCCTGAATTACTAAATCTTCTATTTTTGAAGTTTTACTAAGATTTTATTGTATT
TCCCTGAATTACTAAATCTTCTGTTTTTGAAGTTTTACTAAATACTTCAGAAGCTTTACTAATTAAAAAGGTAATTTA
TAATGTTTTATCACTAACAGTTGATAATAAGCGCTTCCCTATAGCTTCTTAAGATAATAGCTAGAAAACAAAGCTGAT
TTTAATTATTCTTGTAATTTGCTTCAACTTCACTGACAGTCTGTTGTATATTTTCTGCATATGTAATTACATCAGGTTT
TGACTATTTTCTCTGAGGTAAATAATTCCAATTACTTTAACCTGTCATTGTTTCTCTTCTCTCTCATTTCTTCTCTC
TCCCTCCCTCTCTCTTTCTCTTATTATAAGACATATTATTTGTTTAAATTTAATAGCTATTCCAGCAGTTAAATATTCT
TATCATAAATTTCTTGCAATTTTAAATGTGGAATTCCAGTACAAAACCTTTGGTTTAGCAGTAGGTAAAAAGATAGATT
GAAATTTATGTAAACATATTATTCTTCATCAGGCAGGGCTAGCTGACTGACATCTGCTTTGGTTTGGATTCAAATACAG
ATTTAAAGGGAGGCTATACATGGAAATTATCCCCAGGCTCCTCCCCGCTGCTGTTCCCAATTTCTGCAGCAGCCTGCAT
GCTTGATTTTAGGATTCACCAATTTTTTTTGGCCCTTGTGTGTGCTTGGCTTTTGTTTTGATTTACTCTACAATGGCAGG
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GTCAAGGAAATTCATATCTCTGACTAACAGAAAATATTCTTTCACAAATAAAATGTGAAGTGTTCTTAGTTGTCCACT
ATTCTTTTAGGTCCTACATTATTCTATATCTATTTTCAACAAAAGCATAAATTATGGGAACATTGAATTAAATAACAT
GAAGGTCCATACCATGTCCCTGGATCAGAAGTCTCAAATCATAAAAATTGCAGTGGTCCATCAACTGATCTGTAGATT
TGATATAATTTAAGTCAAAATACTGACTTGTTTATTTTGGGGGAACCTTGCCAATCTGGGTCTAAAACTTACACGGAAAA
GCAAAAATTAATAGGCCACTGTAGATTTTCTTACGAGGTAGTGTGAGTTTTTAAATTCATTTATTTATTTTGA
ACATTTAATTTAGGTTTGGGGTATATGTGCAATTTTGTATATAAAGGTAAACTCGTGTACAAAAGCATTGAATTTTATG
CCATGTTATCTATGGTTGAAATATAATAGGGAGCGAGTTTATTTTAAATCATGTGTCTTTTTTAAGATTTGATTTATGC
TGACTTCAGAATGTGACGAGTTTTAGAAATCTGAGTGTAGGAATGAATGTTCTATTGGAAGTATCTAGTTTATTGCTTT
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TGTGCCCCAGGCCGACTGCGGACTGCAGTGGCGCAATCTCGGCTCACTGCAAGCTCCGCTTCCCGGGTTACGCCATT
CTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGTGCCCGCCACTGCGCCCCGGCTAATTTTTTGTATTTTATAGTAG
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TGGTTGATGATGGGAATGATTCTGTAACCTGCATTTATGGTCTAAGAGCTCACTGGTGTATATACCTGGGTAAATTGGG
GATTTTAGGTAGCCAGAATTGGAATAAGATGCACAGTATGAGAAATCCCCCGTCTCTGAGGATGGGGGATGCTGGG
GCCCCGTACCATCAAGTAGCACATCCACAACCTCACTCCATTTACCACATCATCACCATCACCACCACCAACAAGCACT
TGACCTCCATCCTAACTTCAACAATCACCTTCAGTCTGCCTAACTGATGCTATGGAAGAACAACAGAGACTCAGAAGGG
TAAGAGGGTGGTGGTGGCGGGTGGTAGATGATGAAAGATTACTTACTGGTGTGAACCTTCAAGATTTGAGACAAGTCTT
AGTTAATTTGGAAAGTTTATTTTGCCAAGATGGCATTGAGGATGTGCACCCCTAACAGCCTCAGGAAGTCCTGACAATA
TATACCCAAGGTGGTTAGGGCATAGCTTAGTTTTATACATTTTAGAGAGACATGAGACATCAATCAATGTATGTAAGAA
GTGCACTGATTTCAGTCTGGAAAGGCGGGACAACCTTGAAGCAAAGGCAGGAAGACTGGAAGCGGGAACCTTACAGGTCACA
GATAAGTGAGATGAATGGTTGCATTATTTTGAAGTTTCTGATTAGCCTTTTTTAAAGGAGGCAATCAGATATGCATCTATC
TCAGTGAGCAGAGGGGTGACTTTGAATAGAATGGGAGGCAGGTTGGCCCTAAACAGTTCCAGCTTGACTTTTCTTTT
AGCTTAGTGATTTGGGGGCCCCAAGATTTATTTTCTTTTCACTGGGTATAAGGTACATGATTTGAGTGATGGATACC
CTAAAAGCCCTGACTTCACTGCTATGCAATCTATGCATGTAACAAAATTATACTTGTACCCCATACATTTATACAAGTA
AAAATAAATAAATAACAAAAACAAAACAAAATAACAATGGCCCTAGCCATAAAGCAATTTTTAATTTAGCGGGGAGGA
CAGATATGTATCTAACCAGTTATACTGGAAAGTGAGTGAGCTCAATATGCTAAATAGAAATTATTGTGCATCATAAAT
GTTTAAGAGGAGGGAAGGGTAATCTGACTGGAGGGATTGAGGAAGATTAGAGAAGTTGTCAATTGACTTGGCCTTGAGG
CAAAACAGGGGTTTATTTATTCAGCAAATCAATATGTATTGAGTGCCTAGAATTTAGGAATCATTGGAAGGGCATACTG
GGAAGAGATTATATTGAGATCAAAGATCAAAGAGAGAAAAAATGCAAGGAAGTTTTTAGAATGTTGAGTCACAGAAGGG
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ATGAACAGAGCATGATTTATTGGTGGATATGTTGGAGGGGAAGGAAAAATTATACTGAGGAAGACCAATTAGAAAAATTT
CACAAATAGTTTAGGCAAACAATGAGGGCAGACAGACCCAGGATAGTGGCTGTAGAAACAAAGGAACAGATGTGAAGAAT
TTTATACAGATGTGTGTTGAGATTGCACAGTGGTGAAGGCAGGGCTTTGGGGGCACCCATTACTCAAATAATGTACATT
GTATCCAGTGTGAAAGGAAAAATAATCTTGGGGCCCCCAAAACATGAAGCTAAAGGAAAAAGTCAAGTTGAGAGACAGA
AAAGATAGAAGAAAAAGGCAAAGAGAGCTCAAAGGTTATCAGCCTGACAGTGATAAGTGATTAACAGAGGAAGAGTGTC
TGGGTCCATGAAATAGTGATGTTTTGGGCAAATAGAGTCTATGTTGTCTGGGGTGAAGCTAAAAGAATCATTTAGCA

Fig. 9.125

Fig. 9.126

ATAGCTGCCATTATTTGACTGCTTATTATGCACCGTGGAGAGTTCTAAATGCTTTAGCTAATTTAATCTTCATGAGACA
CCTAAGAGATAAATCCTGCTATTATTATTATCATATTTTGGATAGGAGATTAAGGCACAGAGAGGTCGTACAAACAAGTGA
GTGTCAGGATTCAAAGAAGGAGGTTTTGCTCCAGAGGCCTCCTGAATTTAACCCTGTGCTAGAGACACAGGGACATTC
AATGGAGCATTAAATGTCCCGAGCTGGTGACAGGTGAGCTGAGGCTATAGATGTGTTGAGGAGTCAGCATCAGTGGAGG
AGGAAAGAGATGCGGATGGAGGGAGACTCGGTGTCTTTGAAATAGATAAGAGAGAGAGACAATGTAATGGTTTCCACTT
GGCACAGAAAGAAATGCCTTACTAATTAGAGGACCAGGGAGACACAAGAGGAGCCTAAGAAGAATGGAATGAATTAGAA
TCATCACTGTGGAGTCTGTGATAGGGAGTCAACTAAAGTGGAAATAAGTAATTACCAAATAACAATGAGGGGTCAGCAT
CGGCAAAGGAATATGTTTGTAAATAGAGGCTTCGCAGCTGTATCTGGAGCCTGGGAGCGCAGGCTGAGAGGGACATTGAG
ATGTAATGAGGGATGTGACATCTTAGACAGAGAGAGAGCTGGGATGCTCTCAAGAACTTCATTGATATCCCTGTCTGTA
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ATTTCTTCTACAGGAATACTCCTCTGTTATAAGGATTCAATAAATATGGTGACTTTTATTGTTATTTCGAAAGATTATAGG
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AAATTGGAATCTTTGACAAAACACAGCTGGAAGACAGCAGCCAGTAATACAGCAGCCATTAATACAGCTCCATGTCCA
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TTGACTTGAAATTTGACCTCAGCTTTTCATCCAGTATGCTAATGATGGTATAAAATTTACACTCCTTACATAAATAAAT
GTACCCTCAAATGTGTCAATTTCTGCTATTGACAGATGGATCCCATGCTTAAATTTACCACAAATTTCTTATTCTGAGCA
GCTTTGCACAGCACTCTGCATCCACTTTTGTGTTTGGCCACGTTATAGCACTGCTTGTGCCCTGAGCTTGCCTCTCAAG
GTGGCTACAGCGTGACATTTCTCTGAAGAGCGATAGGCAGAATTTAATTGATTTTCAATTTGCTTGGATTCTCAAAGGC
TTCTGTGGCTATGCCTATTACCTAACACTGCTCTCAGGAGGTATAAAGCTGTGTCCAGTTGTTCTTGTGTCATGTCAAT
GAGTGACTTGGTGCCACTTGCAGCAGCCCTTCTTGCAATGACTGCATTCCTGAATCTATTATGACATAGAGACT
CTAGGGACCAATGAGGTTTTGTGTAGGAGGGCTAACTTTTATTTTCCCCTTATAGTTCCTTTGTCTGGATTTTCCCCCT
TTCCTATCTATCTTTATAAAGAGACCTTAAATGAAGGCTACAGCTATAAGATGAACAAATAGCTGGCTATTAAAAATCT
CCAAATTGTCATATAAATGCAAACATGATCCCAATGATTGGTTTGAATAAAATTCAGACTTTACTGATTGGAGGTGGG
CAATTTCCAGTTATAGGCTGACCTTCCACTTCTTCAGAGCTAACCTCTAAACAAGATTAAGCTTATGTCTAGGATGG
GAGAGAGAATGGCTGGAAGAAAAGAAGATGTTTCACTTCTTATGTTGTGGCTTAATGTAATGGCTTAAGAACAATAAT
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GAACCAATAGGCTGTGTGTGTGTATATGTATACATACAGCCTGTATGTTTGTGTGTGTATGTGTGTATACATAAACATA
CAAATGAGGCATGCCTCATTTTATTGTGTTTTGCTTTATGGTGCTTGACAGATATTTTATTTTTTTCACCAATTGAAGGT
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CTATCATAATTGTGTTAGGGTTTCAAAAACCATGCCTATATAAGATTGTGAACCTAAGTGATAGATTGTGTGTTTTCTG
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CAATATTAATAATTAGGCCAGTTAATAATCCTACAATGGCCTCTACATGTTCAAGAGTTTTCAATCTTTTTCTTTAAATT
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AGCAGCTATCCAAGTTGTGAATGCAAAAAAATAAAATAAAAGTTCTTGAAGTTCTGAATTTCTTGAAGCAAAGAAGTTCT
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AGTTTTAGTGGTCTGGACAGAAGGTTCAAACCAGCCACAACATTTCCCTTAAACCAAAGCCTAATCTCAAGCAAGCGCCA
AACTCTCCTTATTTCTTTGAAGGCTGAGAGAGGTGAGGAAGCTGAAGAAGAAAAGTCAGATGCTAGGAGAGGTGGTG
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AGTTACCCAGACCTAGCTAATGCCATTGATCACAGTGGCTACACTAAATAACAGATTTCAATGTAGAGGAAACAGCCTT
ACATTGGAAGAGATGTCACCTAGGACTTTTCATAGTTAGAGTGGAGAAGTCAGTGCCTAGCTTCAAAGGACAGGTTGACT
CTACTCTTAGGGCTAATGCAGCTGGCGACTTTAAATTAAAGCCAGTGCTCATTTAGCATTTCAAAAATCCTAGGGCTCT
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GCAGGTTGTTTCACTTTCCATGTAGTTGTGTGGTTTTGAATGAGTTTCTTAATCTTGAGTTCCAGTTTGACTGTGATGTG
GTCTGAGAGACTGTTATGATTTTCACTTCAATTTGCTGAGGAGTGTCTTACTTCCAATTATGTGATTGATTTTAG
AGTAAGTTGTCTTGTGGCACCAGAAGAATGTATATTCTATTGTTTTTGGGTGGAGAGTTCTGCAGATACCTATCAGGT
TCACTTGACCTAGAGCTGAGTTTCAAGTCTGAAATATCCTTGTAAATTTTCTGTCTTGATTATCTGCCTAATATTGACAG
TAGGGTGTTTAAGTCTACCACTGTTATTGTGTCTAAGTCTCTTGGTAGGTCTCTAAGAACCTTGTCTATGAATCTGGCT
GTTTATGATTTGGGTGCATATATATTAGGATAGTTAGCCCTTCTTGTGTAATTAATCCCTTTACCATTATGTAATGTC
CTTCTTTGTCTCTTTTGTATCTTTGTTGGTTTTAAATCTGTTTTGTGAGAACTAGGATTGCAACCCCTGCTTTTTTCCG
CTTTCTATTTTCTTGGTAAATTATCCTCCATCCCTTTATTTTGAGCCTTTGTGTATCTTTGCACATGAGATGAGTCCCT
TGAATACAGCACACTGATGTGTCTTGATTCTTTATCCAGCTTGCCATTCTGTGTCTTTTAATTGGGGCATTTAACCTCAT
TTACATTTAAAGTTAATATTGTTATGTGTAAATTTGATCCTGTCTCATGATGCTAGCTGGTTATTTTGCAGACTTGTT

Fig. 9.127

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GATGCAGTTGCTATATAGTGTCTTGGTCTTTATATTTTGGTATGGTTTTGCAGTGGTTTTGTAATGGTTTTCTCCTTTAC
ATAGTGCTTCCTTCAGGAGCTCTTGCAAGGCAGGCCTGGTGGTGACTAATTCCCTCATCATTTGCTTGTTTAAAAAGGA
TTTTATTTCTCCTTTGCTTATGAAGTTTAGTTTGGCTGCTTTGAAATTCTGGGTTTGAAATTCTTTTTTTTTTTTAAAGA
ATGTTGAATCTTGGCCCCCAATCTCTTCTAGCTTGTAGGGTTTCTTCTGAAAGGTCTGCTGTTAATCTGATGGACATGA
ACAGACACTTCTCAAAAGTAGACATACATACAGCCAACAGACACATGAGAAAAAGCTCAACATTATTGATCATTACAGA
AATGCAAATCAAAACCACAATGAGATACCATCTCATGCCGGTCAGAATTGCGATTATTAAAAGGTCAAGAAACAACAGA
TACTGTTGAGGCTGTGGAGAAATAGGAACGCTTTTACACTGTGGCTGGGAATGTAAATTAGTTCAACCATTGTGGAAGA
CAGTGTGGCAATTCCTCAAAGACCTAGAACCAGAAATACCATTTGACTCAGCAATCCCATTACTGGGTATATGCCCAA
GGAATATAAATCATTCTATTATAAAAAATACATGCACATGTATGTTTATTGCAGCACTATTACAAATAGCAAAGGCATGG
AATCAACCCAAATGCCACCAATGATAGACTGCATAAAGAAAAATATGGTACATATACACCATGGAACACTATGCAGTCA
TAAAAAAAATGAGATTATATTCTTTGCAGGGACATGGATGGAGCTGGAAGCCATTATTCTCAGCAAACCTAACTCAGGAA
CAGAAAACGAAACACCACATGTTCTCACTTATAAGTGGGAGCTGAACAATGAGAACACATGGACAAAGGGAGGGGAACA
ACACACACTGGGGCCCATTCAGGGGGTGCTAGGGGAGGGAGAGCATCAGGATAAATAGCTAATGCATGTGGGGCTTAAG
TCCTAGGTGATGAGTTGATAGGTGCAACAAACCACCAGGACACACATTTATCTATGTGACAAACCTGCATGTCCTGCAC
GTGTATCCAACAACCTTAAATTAATTAATTAATTAATATGCTAAATCTACTCTCTTTGTGCTCTATAAATGGAACAACAA
AGCCTGTATTACCGTATTTCTGTCTGCAGCATGATATACTGAATATTTTAAGCCCACTATTGAGACCTACTGCTTAGGA
AAAAGAGATTTATTTCAAAATATTACTGCTCATTGACAACGCACCTGATCACTCAAGAACTCTGATGAAGGTATACAAG
AAGATGAATGTTTCTCTCATGCCTGCAAAAACAACATTCATTCTGCAATCAATGGATCTAGGAGTCATTTTGACTTTCA
AGTCATACTATTTAAGAAACACATTTTCATACAGCTATAGCTGCTATAGATAGTGATTTCTCTAAAGGATCTGGGCAAAG
TAAATTTTAAACTTCTGGAAAGTATTTACTACTCTAGATACAATTAAGAACATTTATGACTCTCAGGAAGAGGTCAAG
ATATCAACATTTACCGGCATTTGGAAGAAGTTGATGCTAACCTCATGATGACTTTGAAAGTTTCAAGGTTGCAGTAGA
GGAAGTAACTGCAGAAGTGGCTGAATTGCTGCAATTTTATGATAAACTTGAAGAGATGAGGAGCTACTTCTTATGGGA
GCCAAGAAAGTGATTTCTTGAAATGGAATCTACATCTGATGAAGACGCTGTGAACATTTGTTGAAATAGCAGCAAAGGTT
TTAGAATATAATATAAACTTACTTGATAAAGCAGTGGCAGGGCTTGAGAGAACTGACTTCTATTTTGAAAGAAGATATA
CTGAGGGAAAAATGTTATCAAACAGCACCACATGCTACAGAGAAATCTTTCATGAAAGGAAGAGTCAATGGATGTGGCA
AACTTCATTGTTGTATTATTTTAAGAAATTGACATAGCCACCCCAACCTTCACAACCCCACTCTGATCAGTCATCAGC
TACCACCATTTAGGCAAGACCCTCTGTGACGAAAAAGAGTACAAGTTGCTGAAGGCTCAGGTGATTGTTAGCATTTTCT
AGTATAAAGTATTTTAACTAAAGTGTGTACACTTTTGTAGTTACAATGCCATTACACACCTAATAGACTGCAGGATAGT
GCAATGTAAACATAACTTTTATATTTCACTGAGAAACAAAAAATTCATGTGAGTTACTGTATTACAATATTTGCTTTAT
TGCAGTGGTCTGAACTGTACCTGCTATATCTTTGAGGTATGCCTGTGTGTATGTGTATATATATGTATGTATACACAC
ACACAAACACACACACACACACACACACACAGAGAGAGAGAGAGAACAGAGATTATTTTAAGAATGGCGTGAACC
CAGGAGGCAGAGCTTGCAGTGAGCCGAGATCGTGCCACTGCACTCCAGCCTGGGTGACTGAGTAAGACTCTGTCTCAAA
AAAAAAAAAAAAAAGAATTGGTTCATTTGGTTCATGGAAGTTGGTGAGTCCAAAACGTGCAGGATGGTCCAGCAGACTGAA
GACCCAAGGAGTGGATTTTGCAACTCAAATCCGAAGGCTATCAACTGGCAGATTTCCCTCTTCTTTGAGAACATCAGT
TGTTTTTTGTTATTGTTGCTCTTAAGGCCTTCAATTTCTTTGAGTGAGGCCCATCCACATTATGGAGGGTACTCTATTTTA
CTCATTTACTGATTAATCTCATCTAAAAAGTAGTTTCACGGCAATATAGATATGTTTGACCAATATCCAAGTACCATG
GGCTAGCCAAGCTGACATATAAAGTTAATCATCATACTCTCTGTGTAGTGTCTCATCATTTAGCCCAAAGAAGCTTGGG
CTTCTTTACAGCATAGCAGCTGGCTTCCCAGGGAGAGCAAGTGGGTCTGTGACACCTGGGCTCAGGAGTCCCGGGATAT
CATTACTGTTGCATTCTATTGGTGAAAGAAAGTCACAGGTCTAGCTCATAGCTAAGGGGAAGAAACTACAACCTACTT
CATGGTGTGAGGAATAGCATATGTGGGCAGGGATGAGGGGAATAGTTGGAGACTAGCTATCACAATCTTCCCTCTGGCC
ACAGCACTTCTTGTCCCTTTTCAGAAAGTCTCATCACATTACAGTGCTGGGCCAGCTTCAAGGTCCAAAATCACATAA
TCTAAATTGGGTATATTTAAAGTTGATGTTCCCTCTTTATCTAGATACTTGTACCTAAAATGATACCTTCTGTGTAATC
CCCCTTCCCTGCAACATACATTGATGAGACAGGGATTGTTGTATCGCTAGCAAGAAAGTCAGCCATTCAAAGGAGGGA
AATGAGAGGCACATAGCAATCGGTGCTTCATAAAAATTTCTGAGATCCAGCTGGGAACATGTTACCAATTCCCTCAAATC
TAGGTTTCCCTGAAAAGGATCCTGTTCTCTGGATGTGGATTTCTATTCCATTGTTTTTCAAGTGCCTGTTGACTCTCTCCTC
TAAGTTAACCTTCCCTTTCTGTAAAGAAATGGCCCTCATTGCAACCGGAGTCACTTTGTGAGACTATTTATTCCATTAA
AGGTTTGGAGGCCATAAATCTTCTTTTCATTTTGTCTGTCACTCTCCCTTTTCTTCCAAGGTGGCACAGCTTCTTTTT
AACTTTGTAAATTTTCTAATGTATCCAGTTATATTCCATTCCATTGGGCGAAAGCTATTTCCCAAGTCTCTTTAAGAC
AGACACTTTTCTTCTTTAGACTGAGAGTCAGAATCCTGTGAGTAACGTTTTTAAGAGTTTAATCACCTTGTCTCCTA
GCAGAGTGGAACCTTATAAGGGTTTAAAGAGGTATCTTACTACCACATTCTTGACTTGATATTTACCCTGAGGCCACATT
GTACTAGCAGTACTTGATTTGATCAGAGACCATTTTTTACTCTGAAAACCTTCTGCCATCTGGAGGGGTTGAGAATGAG
AAATAATATTATTTTCTAAGCCAGCAAGTCCTGAGTTGGACTTTGTGGATTAAATAACAGTTTCTTATTTCTGATGGG
TAGGATTTTCTGAGTATGTCTTCTGCTCCATGTGGTATTGATTAGGTAGGTCACTTACGTGGCTACATTTAGTCAATA
GCTGGTCTGGGCTGGATGGTCCAAGAAAGTTTCAGTCATATACCTGGCACCTCTGAGCTTTTCCATATGGCCTCTATGT
GGAATCATCTTGGGCTTCCTAATAGTATGGCTGATTGAGACTGGTCAGACTTTTTTTTTTTTTTTTTTTTTTAAAGTGG
AGTCTTGCTCTGTTGCCAGGCTGGAGTGCAAGTGGCATGATCTCGGCTCACGGCAACCTCCGCCTCCTGGGTTCAGCAA
TTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGACTACAGGTGTGCACCACCATGCCAGCTAATTTTTGTATTTTTTAGC
AGAGATGGGGTTTACCATTGTTGGCCAGGATGGTCTCGGTCTCTTGCAAGGCTGGTCAGATTTTTTATATGGAGGCTGAC
TCCAAGGGAAGGTGTTTTAAAGGAACAGGCCTGCCAATGTGCAAGCACTTATCTAGCATTTGCTTGCAATAATTTTTCT
AATGTGCCAAAGTAAGTCATATGGCAAGGCCAGGATCAACATGGGAGAACTACATGAAGTGGCAGTGCCAAGAGGTAG
CATTTATTATGGGTCACCCATATAACAATCTCTCACACAGAGTAGTTTGGTAGCTGGACAGAAATAAATTTGTATCTT

Fig. 9.128

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CTTGTAACCTTTTACTAGTAGTATAACTTAGGGCAAATTGCTGTGCTTCTGGGATTCTTATTTGCTGCTTGTAATAATAGA
AAACATGACACAGATGCTACTAGATCTTTGTGACAACTGTACACACATATGCATACACACACACAGAGGTACTCACAG
TGGTACTTAACAATGGCTACTATGTTTTAATGGTGTACTTGGCAGACCAGCAGTTAGGTTTTGAATGGACTAACTGTGG
TTTTATCATATCAGGACCAGGTTGTAATCAGAAATCAGCGCTTGCATAGCTCAAGGTGATTAAAGGTAAAAAGGTTAAA
AGTTGAGAAGAAAGGTTGGTGTAGATACTATTAGTTCTTGGCCAAAAGCTGTTGGCCTTGGGAGAGGTGGAAGTTTGA
GATCAGGGATCACAGTTATATGAGAACACACAGTGATGGTGAAGTTTGGAAACCAGAAAATAAGTCAGGATTATACGGGG
TCCAAAAAAGCGGAGTCAGGGCACTCATGAATCTAAACCAGATATAGAAGCTTATGGAGTCAGAGAGCAGAAATCAGGG
ACACAGGCAAGAAATCCAGGAAAACAAGAATTAGATATACCCAGTATGTTGAGACAAATAGCTATGGGGCGCAGAGGCA
AGAGAAAAATTTGGTAGTTTGGAGCCATTGTCTCCAAACATAATTATAGCCAAAAGTAAAACATGTTACAGCATGTGG
GCAGATCTTGAGCCATAGGTGGAAAAAGACATCTGGTTAGAATACGATGGCAGCAAATTTGGTAGCTCCTGACATGCATC
AGAACCCCTTTAAATTTTCATGGGATTCAGGTTGTTCTTCAATTATTTCAATGATCTAATCTTTAACTTCTTGAGAAT
ATGATGTTTTCTTAAAAAGGGGACTCCACACAGAACTGTGAGGCTATGAAGCTGTGATGGCATTGTAGTACTCAC
ACAGTTCAAAAAAATGATGCTATCTCTGCCATCTTTTGTTCATCTGTAGAATAGTCCATTTTTTTTTTGGTTATAGCAGT
CGTCTTCTTGAATGGTTCCTACCCAGAGGGAAATCGGAATTGCCGGTACTCCTATGTCTCAGTTGCAGAAGTCAGGAT
TTTTTCTGAATCATTCTGAGCATCTTATCTTCTCATGCTGATTATTTATTTAGTCTGAAAGCTTATGCTCTCTTATA
AAGTCACATTGTACTTTCGTTTTCCATTGAAAATTTCTGAATAGAATAAAATGAAATCATTAAAGTCATGTTTAAGAAA
AGTAGAGTTTCTTGATTAAAAAGGAGAAATTTAGAGTGGCATTCAAAGAAATACATTAAAGGAAAAAGTAGCATGCA
CTCATGCAGTTCTTGTGGGAGGATAGTTTCTTTCTCTCATTCGGCCCATTAATAAGTTCAATTAATTTCTTTTAA
AGATTTCAATTATGAATGTATCTTAGGCATTGATAGGGAGTAATTTTTAGATCTGACTTATTTTTATATATTATTGTTT
ATTTTTATAGCTTTTCTGAGATTCTTGGCAGTGAGTCAGCACACAGCATTTCTACCTATAGTCCAATTTATATGAGTCC
TACTTTTCATGCACCACTGTGATAGTGATGTCTCGGTTAGTGGCTATCATGGTCAGTAGCCTGAAGATCTATAGCATCTA
CATATTTAGAATGGATTTTTAATGTCTATGAGAGCCTTTATTTCTCCACTACCCGGTTCCTTTGTGGATCCTGAGTCTG
AGGAGAACATAACTGTCAATTAGAACCCTTTTAAAGAAATFGCTGTAATTAAGATTTTTTGATGACCGAGTTATTAATTA
GGAAACAGGGTCTCCATATTTTGCTCTGCTGCTTCTTTATGTCTTAGAGCAAGGGTCAGCATACTTTTACTGTAAATGG
CCTGGTAGTAAATATTTCCAGCTTTGTGGGCCACATCATTTCTGCTACAAATACTAAACTGCCATGTGGTACAACAGCA
GAAAAGGACCTATGCATAGACTGCTGTTTTTAATGTCAATATAGTATCAGTAGAAAAGACAGGGCTTAATAAAAACTTT
GTGTAATATGGTCTCTTTGGCATAAAATTAAGAAATGTATAGAAGTAAACGTTGTACAAAAACTTTTGCAAATGTGTG
TCGCTTTTCCCAGCACTCACGAATAACAGTCATTTTAAACCAGACTTCTCGTTTTTGCCAAGTATTATTTGTTCTCATC
TAGTAACAGTCTAGGACAGGTGTGAGCAAACTACTGCCACAGGCTAAATCCTGCACCTTGTCTTTTCTCTCATGCCAC
ATAGATATTCAAATGAGGAATCTATGCACAGGTCTTTGACCCACCCATCAGCAGAACATGCCCTTTCTGGAGCTCCATT
TCTCACTCTCAATAGTGCTGACCACTTAGGAGGAGCTGTCCATGTTCTTTTGTGTTGAAATTTCTCTATGAAAAATGACAT
CTTTTCTGAAGTGATTGCTGGAATAATGGGGCCTTTTGTGTTTGCAGACCTGTTTTCTCTCAGTGAAGCTGGCAGGGAC
TGGTAAGGAATCAGAATTACAATCTTGTCTTTATTAGCACTGTGTTTGCATCAGTCCTCTCCTGAGAAACAGTGCAGGT
TGAGATAAATTTCTCCTGTGAAGTGATACAATTCAATTTTCATCTCACATATGCATGGCCTTTGTGCCATGCAGAACACA
CTTTTTACTGTTGTTGATGTAAGTGAGATGTTATTAAGAGGTTTTAGTTTTGAGTTTTATTTTAGATAATAACCAATAATTG
AATGCTTTACTAGTGTGTTACTACAAAGGTTTAAAAAATCAATTAACCTATAAAACTGAGTAAATAAAAAATGATTTAG
AACTAAAGATAATCTCAAATTTGCACATTAGATAGCTATCCTATGTTGTAGAAGATATTCAGTCTGCATCATAATATTT
GAAACAAATACAACTTTTACCATAAGACAAGAGCAAAATGCACAAAGAATCGAGTGTCCATGCAATAGGCTTTATGA
ACACAAAGCCTGTGGCCAAAATGAGGCAAGAGGAAGTATGAGGGGTACACAGATGTGATGTGAGTACCACATAACCACC
CCTGGGGAGTGATAAACTTTTTTGTGTATTTGCATTTGACTGATTTTTACTTTGTGAAAACATAATGTGCTGGGGAAAA
TCACATCTGAATCCACAAAATGTCCATGGTATATTGACATGTACCCCTAATTTATCAATTTTGTATGAATGTATTTGCA
TTAGAGAAACATGTTTCAGAATACATTGTGCTTTGCAAGTGTTTGCACCTGTTGCAAACTATGTGCTTTATCTCACTGAA
TCTTCACAATTACTGCATGCGGTGGGCACTATTTTCATGCCTGTTCTACAAATGAGTAGACATATAAAAGTTAAATAAC
TTGCCAGTGGCTTTGAGGAAATACGTAACAGTATATTAGAGATGGGATTGAGTTATTCTAATTTTCAGGAAGAAGAGGA
AAGTGAATTTCTCCCACTTATGTCTTAGGGAGTAGGACATTAATTGTATTTCTTTTAGTTCTTTTTTGCAAATGTTACT
TTCACAGAGGCTTCTCATGGATCCCTATCTAAAATTACTCCCACTTCTATAGACTTTTTGTTTTAAAAAAATAGAGGCCA
CAATGTACATACTCCAAGACCACTGCCATTAGCCACATAACCAAAATTTAAATTATCTCAATTTTCTCCAAAATACTA
GGTCTAACCATAAACAACGTTGAAATGTGAGCTTTTCATCCTTGTCAACATGACTCAGTAAATTAACCAATCAGCT
GCAGACAAATCAGCTTAAACAGTTTTACTTGTCTTAAAGGAATATAAGTTTATGATAGCCAACCACAGCGAAGACAGA
TGCAGTTCCTTAATTATGCTTTATAAGCTGCATTTTAAATGCTGTGAACAGAGCTTCTTACCACTTTTGATTTGAGGTC
CCTGGTTTGCAAACGTGCTTTTGTATGCTCAATAAACTTTAAAAAATTTTTCTAACTTGATCTGATTTTAAACACATTC
TATCTACTTTAGTGCTTTTTTATATCCTAAACACATGACGCTAACATTCTATATAGCTCACTTGTGTTTATACTGTTTATTT
TCTCTTCCCCTACACTAGTGAAATTTCTATGAAGGCAGGAATGTTTGTGATTGTGCACTACTGTATTTCCAGTGCCTAA
TATAGGCACTCGATAAATAATTATTGAATGAATGAATCAATCAATTGATTAACTTAGGAAAAGTTCTTTAATGGATCT
TAAAAAAAATCTTGTGAGCCTTTAGAAAGGAAAGTGATGATCACCTGGAGCTTGTATTAGGTCACTAATAGTAAAGCA
TATCAGATTAAATAACTTACTTTTCCATAAGACCAGTGATCTGAGGAGTGCTGCTCCCCCTGTCTATCTGAATTTCA
ACGTTTGATAAAAACCTTCCATGATGAACCTTATGCTTAAGATGTAGGAAGGCAGTTGATGATACTGTTAGGCAGAGTAA
CAGCTGAATGCCTGTATTCATGAATCAATGTTTAGAGAGAGACCTCTAGTAATTGGCTGAAGGATTTGGTCTTCACATT
TCCTGTTTACCATTTTTCATTGTTAATTTTGAGGGTGGTATTAATGGCATAAGGAGCTAATCCACAGATGAAATAAAAT
GGCAGGAACCACTAATATGGCTGTTAACAATAATCAGCATACAAAATTACTTGACAGACTAAATGGGAGAGGTAAATATA
ATAAGACAAAATACTAAATGAAAATATAAAGTCTAAACACCAGAGGGAAGTAGGAACTAATGTTGTTTGAGGAGCAA

Fig. 9.129

TAGGAGCAAAGGTGAGACATAGTTTAATAACAACCTGTAGGTTTAGGTGGTAGTTACAGAAGCATAAATCAATAATATGA
CATTATCCCCCAAATATTCATTCTTGGGTTTCATGAATAGAAGCTGGGTTTAGAATAGAAAAGGTAATGATTTG
GCTGTAAATTTATGTTGCTTCTCCCATAGCTAGAGCTATGACCCCAATCCTGGGCAATTCACCTTTAAGAAGCTTTGAGAA
GTAGTGTACACCTAAACCTTGCAATCCTCATGGAAGATGAGTCCAAACTGTGTCACGTGAAGAATACCTAAAGCTGTG
GTTCCAACTTATGGCCTTCAGATGTGTTCTAAATGGCCCATATGGGGTTTTTAGAAATTTGAATTTGCGGCAGATGTT
TCAAAATTAGTAGATGTCTCATTAAATATAGATTTCTGGGATTTACTCTTGCTACTGTGAGGGTGTGTGATCCTGTGTT
TGTGTTCTCTCAGACCACTTTACTCATTACTTTAAATTTCTGGTGCTATAGCAATCGGGGTGCTAACACCCGGGGG
GATAAAAAGGACTGGGGAATTCATGGCCTGTAAACAAGGTTGGGGTTGGAGAGATATAAATATATCTGGAGAGCTTT
CATAGGGCAGAGCAATTAGACCCAATCTTTGTGTCCCCAAATGTTAAACACTGGTATTAATAATAAGAAAACATATTT
CCATTAACCATATAGAATAACTATCTTATAGTGAAGTGGAGACTGGCTATTTTATTCATCTTTTGTTCATCAGTACCGAG
AAGAGTTTCTGACAATGTTTGTACTCAGCCACTCTTAACTGAATAAATGTGGCAAAACAGGTTGCGAGTGGTGAATTCC
ATTCCAGGAGCAGCTAGAATATATGCTCTTTGAGAGCAGAGATCTTTGTCAATTTTGTTTACAGCTATATCAGAGCACC
TAAATCAGTGCCTGCCATGTAGTAGGAGCTAATTAAATATTTGTAGATTTTATAATTGGAGATATTCAAACATAGAATC
ATGGTCTACTTGTGAGTGTGTAGGAAAAGTAATTAACAATATAAGTAGAGGGTCATGAAACAAGGGAAATACTGAAG
TAGATGACATTTAAAGTTACTGCCAACCTGAAAGCCTATCAGACAGTGACATAGAATGGGATTATGTATTTTACATTA
AATATTAAGCATAGTAAAGTTAAAGTAGTGTAGGTATGTGTGGTAACAAAACCTGAGAAAGCAGTTTGCCAGTTATTTAT
TTATGGTATGGACAGGAAATTATTGGCTAGATTTTTTAAATGAAAATATCCAATAGTTGAGATTTTTCTGATATTTAACT
ATATCTTATATCTTATGTCCATATTGATGACAGTGATTATTCATTTTCAATCGTTTTGTGTGTCATCTAAGCAGCAGTTA
ATCATTCCCTACGTGATGCAAAATATACTTTTATTCCCTGGTATTTTGTAAATTTAACTTACAAAACACACGTTGCCCTGAGG
TTTGGCATGATTTGTTATTCTATCCTTCAAGAAGACCCATGATAACAAATCAAATCATTGAGAAGTATTCAGTTTTC
ATTGTTTGGGGTGGTAACCTGAAGGTGACGTTTTGAAAAATCATACTATAACTAACCTAGAAAATAATGTTTCAGAGGCCA
ACATGTGGCAAAGGAACACATAGTTAAGTATATAGTTTTTAAATAGAAATTCTATGCACCTTTTATTGTGAAGGTCTAGC
ATTTTAAATATTTTCAAATCCAGAAAGATGAAAAATTTCTTTATAAACTTGAAGGGAAAATAAAAAGTCTAAAGCCCTG
AATTTGAGTTCTAGAAATGAATGTATTTAAATGCAGTCTGAGGTGCTGTATGCTAACTATGTAAAGCAATAGCTTAT
TAAATTCCTTGAAACTTGCTGTTTATTGAAATTGGATATGGTAATAGATTTTTGTAAATGTTATTTTATAAGGGATA
ACATTTTCTCCTGCTTAGTTTCAGGAAAAAAGAAAAGAAGCACTTAATTTTTTAAAGGTTAATGGATACCTAATGGTTG
GATTAGACCTATGCATCATCATGTTTACAGCACAGAACAATGGGAAAAATTTCTATTATGGAGCAGTGCCACATAACCT
TAATATCCTGTATCATCATTATCATCATTATTATGTGTTGAGTATACAAGCGTCTATCTCTGAGGAATACTTAGCAATT
TATGAATTTGTTTTAATTAGTTTTCTCAGCCTTCTCATAGGGCATGGGGGACATATTGCTTAACCTAGTTTTAGAGCC
AGCCTTAGGTAAAATGGTGATGCAAACTGTTTTACTTTGGTGACTCTTCTGCTTGCAGGGATACCTCCTTTTTGCAC
AGCACCTTATTTCTCCTTCTGTAACTAAAGGAGAAAATTATTATACATTTTTTATACTTAAAAATACGTAGTTGAGAAT
TTCAAATATATGCAAAAGTACAGAGAATAGTAATTGGTTGGCGCGAAAGTAATTGTGTTTTTTTGCCATTTAATGGCAA
AAACCACAATTGCTTTTCGCCAACCTAATATAATGCATCTTAAATACTCATCACCCAGTTTTTAAACAATTCTCGATATAC
AACTAGTTCTTGTTCATCTGTACCTTTACCCATATTTCTTCTCTACAATTATTTTCAAATTCATAACATAATATTT
CTTCATCTGTAAGTCCTTTAGTATGTATCTAAAAATAGTCTTTTTTAAAAATTACCTCAATAACATTAAGATGCTTGTA
TTTTCTTAAAAAATTTCTTAATAATTTCTTATTATCAAATGTAGCTGATTTTTTAAACAATTTTATTAAATAATTTTAA
TTTTTGGTATGAAATGGTAGCACTAAGGTTAACAGATCGAAATCATTATGTGTCTGAATTTCTCTTTTACACTATAAAT
TTCTCTTTCTATCCTTTCTCTCTGCACCTTTTAAAAAACTGGGTCATACATCTTACAGAGTTTCTCATAGGTTCAA
TTTTGCTGATTGTATCTCTATTGTGTTGTTCAACCTGTCCTTTGAATTTTCTATAATTATTAGTTAAATCTAAAGGCTT
GATAAATGACTAATGTTTTATTATTTTGGAGAGGATAAGCATATTTTGTTTTTTGTTCCTCCAGCTTTATTAAGGTATAA
ATAATAAATAAAAAATTGTATATATTTGAGGTAGACATGTGATGACTTGATATAGGTATACATTGTRTAATGATTACCAC
AATAAAATTAATCAATGCATCCATCCCTATAAATGGTTACCTTATTCTATGTATGTATGTGTGTATATGTGTGTGTG
TGTGTGGTAAGGATTAAGTATTAAGGATACTTAACATTTACTCTCTTAGCAAACCTTAAACAATACAGTATTATTATCTA
TTGTCACCAAGCTGCAATATTAAATCCCCATAATGTATTTGTCTTATAACTGAAAGTTAGTATGCTGTGTCCAACATTA
CCCCATTTCTACCATGTCCCAGATTCTGGCAACCACCATTTCTACTCTCTGTTTCTATGAGTTCAACTATGTTAGATTAC
ACATATAAGCAAGATCATAACAGCATCTGGCTGTCTGTGTCTGGTTTATTTTCTAGTTAGCATAATGTCTCCAGGTTTATC
CATGTTTTCTCAAATGGCAGGATTTGTTTTTCTTTTTACTGTTGAATAATATTCCATTGTATATACACCACAGTTTCTC
TATTAATTCATTTGTCAATGGACACTTAGGTTGGTTCCATACCTTTGGCTATTGTCAATAATGCTGCAATGAAGACGACA
AGTGCAGACATCTCTTCAGCATGCTCCTTTCAATTTCTTTGGATATATACCCAAACATTAGAGTGCTGGATCATATGGT
ATCTCTATTTTTTAATTTTTTGGAGGAATCTTCATACTGTTTTCCATAATGGAGTTACCAATTTACGTTTTTACCAGCAGTG
CTACAAGCGTTCCCAATTCTCCACATAGTCACCAACACTTTTTATGACCATCCTAATGGGTGTGAGAAGATACCGCATT
GTGGTTTTGATGTACATTTCCCTGGTGATTAATGATGGTGAACACCATTTTTATATACCTTTTGGCCATTTGTATGTCAT
CTTTGGAGAAATGTCTATTCAAGTCCCTTGCCTATTTTTTAAATGGGTTATTGGAGTGTTTGCTATTGAGTTGTAGGAT
TTTTTTGTGTATTTTGAACATATCAGGTACACTCAGCCCTTTGTATCTGTGGGTATGCATCAAGGGATTCAATTATCT
GAGAATCAAAAATATTTGAAAAACATAATAAAACAATAAAAATAAATTAAGTAATATAGTATAATTATTACATAG
TGTTTACATAGTATTAGTATTATAAATAACATAGAGATGATTTAAAGTATGCAGGAGGATGTATGTAGGTATATATGCAA
ATACTATGCTATTTTATATGCAGGACTTGAGCATCTGAAGATTTTGGTTTCTGCAGAGGGTGGGAAAGGTTGAACCAAT
CCCCCATGGATACCAAGGTAAACTATGTATGGTTTGCAAATATTTCTATCATCCATATGTAGAAAATATTTCAATTT
GCTGATTGTTTTCTTTGCTGTGCAAAAGCTTTTAAAGTCTGATGTAGTCCCATTTTATTTTTTATTTTGTGCTACACTT
TTGGTATCATATTAAATAATAATCACCAAGACAGTGTCAAGAAGCTTTCCACCTGTTTTCTTCCAGGAGTTTTGTGG
TTTAATGTTTTAATTTCTTTTAAAGCACATTTTTTGTATTGTATGTAATAAATATGGATACAATTTCAATCTTTTGCATA

Fig. 9.130

TGAACATCCAGTTTTTCCCAGGAACATTTATTGAAGAGACTATTATTTCCCCATTGTATATTATTCATGCCCTTGTCAAA
GACTAACTGATTATATATATGCAGGGTTTATTTCTGGGATCTCTATTCTGTTCCATTGTTCTGTGTGTCTGTTTTTATGCC
AGTACCTCACTGTTTTGATTACTATAGCTTTGTAATATAGTTTGGAATTAGGGAGTATGACGCTTCCAACCTTGTCTT
CTTCTTCTTCTTTTTTTTTTTTTTTTTTTTTTTAGATGGAGTCTCGCTCTGTTGCCCAGGCTGGAGTGCAGTGGCATGATCT
TGGCTCACTGCAACCTCTGCCTCCCAGGTTCAAGCAATTCTCTGCCTCAGCCTCTCTAGTAGCTGGGATTACAAGCACC
TGCCACCACGTTCCGGCTAATTTTTGTATTTTTTAGTAGAGATAGGATTTACCATCTTGGCCAGGCTGGTCTTGAACCTCC
TGACCTCATGATCCACCCACCTAGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGTCACCACGCCCAGCCATTCTTCT
TTCTTAGGACTGCTTTGGGTATTTCAGGGTCTTTTTTGGTTCATATGAATTTTACAATGGTTTTCTCTATTCTGTGAA
AAATTCATTGAAATTTTGATAAGGATTGCATTTAATATGTAGATCCCATATGGACATTTTAAACAATATGAAATCTTTC
AATCCATAAATATGGGATATATTTTCATTTTCATTAATGTCTTTAAGTGAGAGATCTTTCACCTCCTTGGGTAAATTC
ATTCCTAAGTATTTTGTCTTTTAGATATTATTAATGGGATTGTTTTCTTAATTTATTTTACAGATAGTTTATTGTT
AGTATATAGAAGTGCAACTAGTTGCTGTATGTTGATTTTATATCTTGTAATTTGCTGAATTATTTCTTAATTATGAC
AGTTTTTTTAGTGAAGTCTTTAGAGTTTTTTTTTAAATATAGAAAATGTCATCTGCAAATCTAATTTGGATGCCTTTTAT
TTCTTTTTCTTGCCCTAATTGCTCTGGCTAGGGCTTTCATTACTGTGTGKAACACAAATGGCAAGAGTAAGCATTCTTGT
TTTGTTCAGATTTTAGAGAAGCAGCTTTCAGTTTTCCACCATTAGTGTGATGTTAGCTGTGGGCTTCTCAAATATGG
CGTTTTATTGTCTTGAGGTTCAATCCTTCTATTCCCTAATTTGTTGAGAGTTTTTGTGATGAAAGGATGTTGGATTTTGTG
AAACACTGTTTTCTGCCTCTATGGAGATGATTATAGATCATATGATCTTTATCTTTCATTGTGTTCATTCCGTATATTAC
ATTTTTTGATTTGTGTATATTGGATCATACTTGCAAATCTAGGATAAATCCACTTAATCATGGTGAATTATATTTTTTA
ATGTATTGTCAAATTCAGTTTGCTAGTATTTTGTTTAGGACTTTTGCATTTATGTTTACCAGGGATACTGCCCTGTAAT
TTTTTTTTCTTATAGGGTCCCTTATCTGATTCTGGTCTGTTGGTAATGCTGGCCTCATAAAATGAATTTGGAAGTGTTCCC
TTCTCTTCACTTTTTTTTGAAGAGTTTGAGAAGCATTGTATTAATTCCTTAAAATGCTTGCTACAATTCACCCTTGTTG
CCACATGGTCCCTGTGCTTTTCTTTGTTAGGAGTTTTTTTTTTTTAATTATGGATTCAATCTTCTTACTTGCTATTGGTCT
GTTGAGGTTTCATGTAATATATTTCCCTATCCCTTTGCTTTTCCAGCCTATGTGTATCCTTAAGGCTTAAGTGAGTCTCTTA
TTGTCCCAATATGAGACAGCATTTTGTTGGATTTTTTTTTTAATTCATTTCAGCCACTTTGTGTCTTTTGATTGAATAATTT
AATCCATTTATATCCAAGATTATTATTTATAGGTAAGGAGTTATTACTGCCATTTAAAAAATTGCTTTCTGATGGGTTT
GTGGTTCTTGTGCTTTTTTTTTTGTGTTGCTGTCTTGTGATTTGTTGATATTTTGTAGTGGTGTGCTTTGATTCTTGT
GTCTTTTCTGGATCTATTAGAGTTTTTTTTTTTTTTTTTCTTTGTAATTACTATGGGACTTTCAAAAAACATCTTGTAGC
ATCTTTTAATTATAATATTTTATTTTAAGCTGATGACAAGTTAATTTCAATCACATACAAAACTCTACACTTTTACTT
TCCCCCGTCATTTTGTGCTACTGATGTATGCTTTGTCATCTTTTTGTATTGCATATCCATTAAACAAATTACTGTAAC
TGGTTTTATTTTTTAATTATTTTACCTTTTAACTTTTATTCTAAATTTAGAAATAATTTATACATCACCATTACCACATGC
AAGTATACAGAATTTAATTATGTATTTACCTTTTCCAGTGAGTTGTATACTTATATATGTATTTATGTTGTTATTTAGC
AGCTTTTCATTCCAACCTTGAAGAATCACATTAGTATTTCTTATAAGGCAGGTCCTTGTGGTAATGAACATCCTCAGTTTT
TGTTTGTCTGGGAATGTCTTTACCTCTCCATTTCTGAAGGCTAACTTTGCAGGGTACAGTATTCTTATTTGACAGGCTT
TTTTCTTTCAATGTTTTTGAATATATTATCCATATTTATTTATTTGTTATCCATGTAGGCTTGCATTGGTGTCTGTGAAT
TTGAAGAAGCAAACAGTCTTTTCGGTCTTTACAGGCTGATTTTAAACAACCTTCTCTTCCACCAATGGCAGACCTGTTATT
AGGTATGCAGATAGGCGTGGTTCCCTCTGGGTTTCTGGAGGACTCCCCCTGGCTCTCTGAGTATGTCTATGGGTAGGGA
GAACCGTCCCCAGATCAACATGAGAGAGCTTGGAACTAAGTCACTGCTGCTTCAGGGTCCACATCTGAGAGGACTTGCC
TCCAGGGAGTTGGATGGGCATACATCTCTGGGGACAAGATTGACCTTGGGCCAAGTCTAAGTGGGAATGGAGCAAAGTT
ATAGGGCCATCTCAGAATCTGCTGTGGAACCAAGTTTGGCAAGCCTGTCTCCGAGAACATGGATGGGAATGAGTCCCTGA
CAGGTCGCTAAGTGGAGACAGGACTGCTCTCAGGCCACAGTTTCAGTTTTAGGATATCATAGTTAGCAGACAGCTCTAT
ACAAAACCTCTACTCCTTTATATCTCTGCCATTTATGTTATCAATGTCACATATTACAGCTTTTTTATATTTTGCATTTG
TTAACATAGTTTTTATATTTATATTTTATAATTAAGTTCTATACCAGAGTTAAAAATAATTTATGCACCACCATTGCAAT
ATTATACGATTTTTTATTTGTCTAAATATTTACCTTTTCTTTTGTGTTGCTGTCTAGCATACTTTGTCTCAACTCAAAGG
ACTCCTTTTAGCAATTCCTGTAAGTCAAGTCTAGTGCTAAAGAACTACCTTAGCCATTGTTTATCTTTATTTCTCCTCC
ATTTTCAAAATACAGTTTTTGCCAGAAAGAGTATGCTTGGTTGACAGACTTTTTCTTTTCCAGCACTTTGAATATATTGTCC
CACTCCCTTCTGGCCTATAATGTTTCTCCTGAGAAATCTGCTAATGTCATTGAAGGTCGCTTGCACATGATGAGCCATT
TTCTCTTCTGCTTTTCCAGGATTCCTTTTTTGTCTTTGATATTTGACAGTTTGATTATAATTTATGTTGGTATGTTGGTT
TTTCCACGTTTGAGTTTTATCAGTGTCTTGAATTTGGTTGTTTCAATTCCTTCTCAGATTTGGCAAGTATCTAGCCATT
GTTTTTTTTCAAATAAGCTTTCTGTCTTCTCTCTTTCTATCATCTCCTTCTCAGATTTGCATTATGTGTATGTTGGAAC
ACTTGATGGTGTCCCATATGTTTCTTAGTCTTTCTTCACTTTTCTTCACTTTCTTTTCTTCTTTTCTTCTTTGTATCTCTGGC
TTCATAATTTTAAAGTGGCCTCTCTTTGAGTTCAATTTATTTCTTCTGCTTGATGGAGTCATCTGTTGAAATGCTCTA
GTGAACCTTTTCAACTCAATTAATGATTCATCTCCATGAATGTTTGTGTTTTCTTTTTGAAGTTTGTGATATTCTCATTT
TGTTCCCTGCATCATTTTCTTAATTTCAATTTAGTTTTCTGTCTGTGCTCTCTTCTAGCACATTGAACTTCTTTAATATTA
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GTGTCATGTTTCCCTGTTTCTTTGTGTGCTAGTTATTTTTTTTTCTGTGATTTGTGCATTTGAAAAAGCAGCCACCTCTT
CAGTCTTTATAAATTGGCTTCATACAGGGGAAGACTTTTACCAATTAGCCACATAGAGATTTTGGGAGCCTCTCCAGT
TTTGTGTTTTGTTTTTTTTTTTTTTTTTACCACGGGATACATCTTCTTTGAATTTGTGTGTGTTCTTTCCAGTTAGAGTGAT
TTGCTTCTTTTTCTTCCAAGAGCTTTTAAATCTCCTTCTCCGTCTGGTGTCTGTCTGTACCACTGCAGGTTCTGTGGCAT
TGCAATAAGCCACTGAGTTACCTTTTGCTCTCTGCAGACCCAGGCATCCAAAGTATGTAGATTCCATTAGTGCTCTGA
ATCAGGCAAGACAGAAACCAGTCTCTCAGACAACACTCTGAAAAACCAGAACATTGGACATACATTCCACTCTTCTCTT
TCTCTCCTAAGGGAGAAGTCATAAGGTGTGATTTATTTTTCTCTGATGACACCAAGCTGTGCTGGCTTGGAGGAAGGGCT

Fig. 9.131

GTCATGGTTGACGTGAAATGCCTTTTCTCATCTGTTTCAATGAGACTATTATTTTCTTTAAGTTTGCCTCAGGCACTGC
AACTTCTTGACTGGTTTCTAGACTTTTTCATAAAACTTTGGATCATATATTATTGCTAAGTTGGCGTCTCTGGTGGAG
AATTAGCTCTCTCGCTGATGAACTCTACAGAATACTTTTACAGTATTTAATATTGCTCTTATTTTCATCTCACACGTAC
ATAGTCACTTTCCTAACACTTATATAAATTGTTCTCTTCCCAATAGCTAGTAGTGTTCCTGTAACACATATTGACACAT
ACTAAGTTCTTATATAAGAAAATATTTAAGGGAATTTTGTATTTTCCACTAAACTTTCTGTTGATTATTTGAAGGTAC
CACACTTTTGGTTGTTATTGTGGCTTCATGGTAAAATTTTATATGTATTAGAGCAAGACCGTAACCATGTTATATTTTA
TAACATGGCCTAATTCCTCCAAAAGTAAAATAAAAAACTAAAAAAGAGATGATTTTCTTGCAATAACATTAACTTAG
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TATTTCTTTCTGAGCTTATTTTAAACATTTTGGTTTGTGTTTGTATATAAATGAGATGTTGGTGTGGTAATTTTAAC
AACTGGTCTTTGATTTTATAAATAAAAGCCTTTTGCATATTTTACTTTAAACTGGACAACTTACCTGACTATTCTTA
TTTTCAATTATTTTTCAGTTGATTTTCTGGGTTTCTACATTAACTAGGTTGTAATCAAATAAGGATAAAGTTGCCT
TCACTTTCTTATAGTTTATATGCCTCCTATTTCTATTCCTCACATTGGGTTACTGAATTTGTTAGAGGTTTTTGAAAA
TGGGGGCAATGTTACATTTGGGCTAAACTATGAACATTGATATCAGCTAAACCTGGTCTCAAACTCTGAAGGATCCT
CAGTTAGAACTGAGAGGTCTTAAAGACGAGGTGGCTAGAGGGAGGTCTGCACACACATTGCCATGAGTCCTCAGACCA
CATGAGACATTCTTTTTTGTCTGAAGCAAGGCTGGAGCCAGGCCTAGATGGCTGTGGAAAGCTCCCATGGAACCAGCC
TGCTGTTCTCATTCCAGGCAAACTAACCTCTGTACAGTTGATAGTGTCACTATGGACAACTAACCTGTTTAAGCTTCA
TTTCCAAATCTGTATCATGGGAATAGTAACAGTTTTTTGTACAGTTTTTATAAGAATCAAGTAGCTACCTATACAATAT
ATGTAATACAATAAATATTACTAAGTTTCTATTATATCCTGATCTTAAGCATGAACCTATGCTTTACCAGTAGCTATA
ATGTTAAATATAAGTTTAATATAAACATTTCTTTTGATCCATAGTTTGTGTTATCAAGGGGATATTTTGTAGTTATCTGAA
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CCTCCACAGCCCAATATTTGTTAGTTTATTCTAGATTTACATACCCAGTTTATCATTATAAATATATCATCACTTAAA
TGATGGTTTTAGGTTTTTCTTTCATGTTTATACACATAGTTTTGATTGGTATTGGTATTTAAATAAACTTGAGAATGA
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TGAGAGAGTAACTGGCTGAGTTTTACCTATACTTACTGTAAGACATAATTTTGTAGCTGCTAAAAAACAATGTTAGTT
TCTTTTCTACACTTGGGATTCTATTACTGTATGTAAGTGTGAAGGAAATAATTGACAGAACTCATGTTTTTCTTATTT
TTTTTCTTCTCTACTAACAAAATAGAATTGACTTTATTTTTTAAAGTCAGCCAGGAAATGTACAATGCTATTTACAA
AAGGTTAATTGAATCCTAGAGTTGCAGATCGAGTTTCTTGTGATGAGAATATACAGTAGAATATTACTGAATGAGAGA
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GACTTGTGTTGTTTCTGTAAAGTTATTTGGGCAATAATAGCATTTCAGTTTTAAGATAAAACATACATAATAAGTAATGA
GACAAGAGGAAAAACATTTGGCATAAGGCTTAGAGAAATTTAAGGCTCATTTTCCATTATTGCTTTAGTGCAATTATTAA
CAATTTTCATGGGGACATGAAAGACTAATATATGGAGAAAATGTCTGCTATTGGCATATAATATGTTAAGATTGCAAATA
TGACTTTTAATCCACTGTGATAATAGCCTATTAAATTTGTATCCTAGTCTTGAAGAGTCACTAACATTATCTTGTGTTAT
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GAATGTGGCATTGGAAGTGATAAAACAAAACACATCCTCTGCCTATGACTGTCTTCTTCTGACCTATCTGACTATTT
AACATGCTGACCTTTACAATGCTCTGCATAGTGTAGGCTCACTGAAAGCACTGAGTGTCTGAAATCCTTCATTCTTATT
TAAACAAAATAGGTGAAAATACCTAATCATGATGTCTATATATTTGAAAGAATATTTGAACATGGGGAGAGACCCAAC
CAGTTTGAGTGGCTGAAGTTTGGAAATTTTATAGAAATTCCTATTATGTGTTTCACTTTATGCCTGACTTATAATA
ATAGTTACAAATACTACATTGCTAGTAACACCAATATCTAATACCATTTCTCTTTCTTAATTTCTTGCAGTTGAAGAA
ATGGAGATGAGTATGTCTATATGCTTCACTCTTTCCTCAGCAAGATTTATTTAGCATATGTTTTGTTCCAGGCACTGGC
TAAGTGCTAGGGGTACAAAAGTTGTTTTGTCTCCCTTCTTCCAGATGTGTCCATGTGTCCAGGCTAACAGAGTGAGTA
GTCACACAGGCAAATCATTTCTCTACAGTTGTGAGGCTGAAGGCTGACTTGGTGTGATAAATAAAAAGGCAGAAGATAT
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ACACCTGTAATCCCAGCACTTTGGGAGGCCGAGGCGGTGGATCACCTGAGGTCAAGAGTTTGAGACCAGCCTGGCCAAC
ATGATGAAACTCCGTCTCTACTAAAAATACAAAAAATTAGCTGGGTGTGGTGGCAGGTGCCTGTAATCCCAGCTACTTG
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AGAAGCAGATAGACAGAAAGCAAAAATAAATGTCCAATTTATGGGGTGATCTACATCTTTTTTTTGTCTGCAACAGTCCA
GGTGTCTCTTTTTTGTGATGTACAACTGGCTAGCACTTTCTTTCACTTGCAAAAGTGTTATGTACTCCAGTAAACC
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AAATGATTAATTGGGAAATTAATAGGAATGGAAACACAATTAAGATAGGTGAACTCCTTTAGAGATATACACATAAAA
TAGTACTACCTGTAACAAGAAAATGCTAAATTTTTGTGGATTACCCCAACAGAAGATTATCTGTGGCTCGTGTACCCCA
AATTAAGTGTTTCTGATTGGCAAGTGCTCTCATGTAAATAATGATTAAGAAACCATGGCTTCTTCCATCTTGTGACTC
TACCATCTTCAACACATGGTGAGGGGCTACAGAGGGGAAGAGAAATAGGGGGTTGTACTCACAAAATTTTTATGTGGCA
GGCCTGAAAGTGGCACATACACAGACCCATACACATTTCTTAACTTATAATCAATGAGCATGTAATGTAACACTCACT
GACATCTCTATAGCTACCTGTAAGGGAGTCTGGAGATGTGGTCTCACTGCATGCCAAGAAAGAAACATTTCTTCTGCA
ATGCCTATATATGTATATGTATAAATATATATATGTATATACATGTTTACATATGTATGTATATATATGTGTGTATA

Fig. 9.132

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CATATACTGTATGTGTGTATATGTATATATACACACACACACATCTTAAGGGGATTTTCTTATAGAAATTTTATTTTC
TCCCCCTCTCTATATATATATGAAATAAATATCAGAGACTGTGAGATTATTAAGAACCACAGAAATTTATTTAACCT
AGTAATGTGCATGCCTGTGTTTTAAAGGCCATTAAGTAGTTTATTGTCTACTTTTGAAAGAAAAAAATTAACAAAGA
AATTAATTAATATCTTGGATAAAAACTGAAACAAACAAAAAGGAAAATAATAATAAATTTAACATTCATCCATTG
TAAGTGTCAATTATTATGGAATTGCTTACTTTAGGCATGTATTTTTCAGAAAGAATTTTATGCAGTAAAAAGCATAACATT
AAAGTTATTTGTCTGTCAATTATTCCATATAAGTTGGCTAATTTTTGAATATAATTTTATTTTACTACAAGGTAGCAGG
TTCATATTACAGTTATTTCAATATGTGAGCATTCTTTTTATTTTGATATTGCATATCTTGAAGCCGAATATTTTCTAA
GTCCCATCAATAGCAAGGTGAATGTTGTATCATTTTTATTATTATGACTTTTTATTATACTTTTTAAAATTGAAAAACAAGA
CATCTTTATTATAGGGAATACAGAAGCTCATAATTAATAAAGATACTAAAACAAAATTATTACCAATATTGCACAGGAG
AGCCATTTTAGAGAATTAGGATACATCCTTAAACAATTGTTTTATGAATATAATATGTGACTTTTTTGTGTTTTGTTTTGA
TATTATTGTTTTTATAAAAAATGGGAGCACTCTATAATCTGTTCTTCCCACTCAACATTGTGTCATTAAACATGATTACGTA
TTCTTCTACATTATTTTAAACAGCCTCATAGTATTTTCATCATATGATGTATCAACATTTACTAAGCCAATCCTTCAGTAC
AGCAATTAGATTACATCCAGTTATTTGCTATTATAAATGAGCTGCAGTGAACATCTTTGTGCTGGATATTTGCCAAAC
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TATTCAGACTCATAAGCCCTACTCATAGAGATTGTGTTTAGGAAGATCTGGAATAAGACCCGGGAGTCTGAATTTTAGC
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TATCTTTTTTCCATCTTCTCCCTGCCATAATGCCATTTTCTAACAATAACATAAGATACTTATTGCTCTGGTGATTAGT
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TAACTGATACACATTGATCTAGAGTGTGAAAAAGCCTCTTATACTGTTTTTGAATGGAAAATGTTAGAATATAGCCCT
CTAGTGCTTTTATCATTTTTTATTGTAAAGATAAAAGTATTTATAGAAAGTGGGTTTTAACTAACAGAGTATAAGCATGAG
TGTAAACTTCATTTTTTAGTAGAGATAATTATCTCAAAAAAGTTACGTCTTGAGGCAGTTTTATCTAAAAAAGAATGTCA
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AGGATACATGTATATGTGAAAGGGAGTTTGTAAAGGAGTATTGACTCACACGATCACAAGGTGAAGTCCCACAATAGGC
GGTCTGTAAGCTGAGGAGCAAGGAAGCCAATTCGAGTCCCAAACTTCAAAAGTAAGTAAAGTGGCAGTGCAAGGATCC
AAAAACCCATCACTCTGTAGAAGCATGTTGTAACATTTGATGATTCTATAATTCTCATCACACTCAAAAGTAGGAAAGC
TGTTTTTCAGTCTGTGGCTCAAGGCCCAAGAGTCCCTGGAAAACCACTTGTGTAGGTCCAAGAGTCCACCTGTTTAAGAA
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GTATCCTTCAATCCAATCAAGTTGACACTCAGTATTAACCATCACAACACATAAGTCATTAACCTTAATCATGACCTGTT
TCCAATCCAGGAATGAAGTTCTGTTCTTAGAGGAACTTCTAAGTACATAAGTACACAGAGGCTCACTTTCCCAATCTA
AAAGCTAAGACTGTTCCAAGTGCATATGCTTTCTGTTTTTTTCTATATTTGTTATCTCACAGAAACATTGATTTGCCAT
TGGTTCTTGCAAAATCAGTATTTTATAACTGGAGGTGGTGTAAAGTTTTATTTTGTATAACCTTTTGATTTTATCCCTA
TGGAGACTAAAGACCAGCAAGGTTAAGACCAGCAGATCTGGAACAGGTATTATAGAAATTTGTATATTATGTTGTTGC
TAGCAAATGGAATTATTGCAAGGAGATGATGGAAAAGATAGTATTTCTGATATCTCCAAGCCATTGTTTTTCAAATGT
GTTGTTCAAACCCCTCCGCATTGGGATCAGAGGTGGTGCATACTAAAATTCAGACTCCAGGTCTTATCCAGATCTGCT
GGTCTTAGCCTTGCTTGTCTTTAGTCTCTGCAGACAAAAAATCAAGACATTATACAAAATAAACCTTAAGACCCCTTCC
AGTTATGAAGTACTCTGATTTTGTAAACAACCAATGGCACCTCTTAAGACCACCTGAGTAATTAAGTGGCAGAGCAGGGGA
TAAAATTTCTATCCTCTCATTTAATGACAGATTGTCATTTCATTTCTAATGTTTATGTAGGGCTGCTCTCATCTCAGTTC
TGTTTAATTTTTTCAAATGAAAATGTGAAGCTTATTGAGTTATGATCAATTAGTGTGTGTTCTGTTATTTTCTACTAATA
CTCACTCTGCCAGTGACTATACTTGTAAAGGTAGTGCAGCATGGTGGCTTTTAGCTCCAGTTTTAACTAGAAGAGATTT
GGGTTTGGATCCTGACATGTAGGGTGATCAATCATCCTGGTTTCCTTGGGACTGTGGGGTTTCCCTGGACATAGGACTT
TGAGTGCTAAAATCAAAGGTCCCAGGCGAAGTAAGATGATTAATCACCCCACTAGCCATGTGGCTCTGTCATGATAGG
TCTCTGGGTTTTTTAATGGGTAAAATTGGAAGCAAGAAGAAGTAACCTCATAGGATTTTAGTAAAAATTAATGAGAC
AGTATATGTAAACACGGCACTACTTAATAAAGACTCAGTAAGTATTAGTCATTATTATTGATGTTTTATAAAGCAAGT
AGATTCCTCAAAGCACAGATTCTCAGAATTTAGGATTTGAAATAAATGAACCTTTCTTTTTAGCTTTGACCTCTTTTTTT
TTGTTTGTTTTTTAAATCTTCTCAGTACCCTTCTGTCTGCTTGTCTATACAACCTGGACAATACAACCCAAGGTTGGATA
GAGTCCACCATCTCACATCCTTTTACCTTTTTTCAAGATATAAATCAGGAAGACAAAAATTTTCTATTTGGGAATCAAT
GTGAAGAGTGAGCATCAAAGACAGCATAGAGCTATGGAGAAAGGGGAAATTAGCCTGGATTATAATAACAGCTGCCATT
TTGTTGAGTGCAAACCTCTTTGCCAGATATTGGTGTCTTCAAATACTTTTATATATTTGGTCCACACAATGACTCCGAGAGG
TAAATATTATTATCTCAGTCTTCAGATGGAGAAACAACCTCAATGAGCCTAAGTGACTTGCCCAAAGATAGTTGTATAG
TAAGGAGTGTAATAAATGAGGTCTGGATGTTCCAAGCTCGATGAATGTCCAACCTTTCAGCCAGCTGGTTAATTTTGAGA
AATATAAACTTCTTACCTGTAAAACCTCAGAAGGGAAGGAAATTGCCATATATGAGTGCTAATCCCAATGCAAGTAATG
AACTATGCACATCATATGCATGATCTCATTTAATTCTTGTATGTAAATATTAGTCCCATTTTTTATAATAAAGAAACAGA
AGCCTTAGAGAGACTAAGCAACACACTATCAATACAAATAGTAAAGTGGCAGTGCAAAGATTCAAAAACCCAACACTCT
ATAGAAGCATGTTATAACATTTGGTGATTCTATAATTCTCAGTATGTGGATATAATAAATATTGCTAGAGTAGACTTTC
CAGCCATTTGGAAATAATTATTGCTCATGTGATAGTATATTTTTCTCAACTAGAATCAAAATATTAACCTTTTGACCTGG
GTTTACCTTTGCATGTGGATAACATGGATCAAATTTGCAAAATCTGCTGAGCTGTGACTTAAATACAGTTCTACTGGG
TCCTCAAGTTTCTTTTAAAGTCTTTTTTTTTTCTTACTACCCTCTCAGTCTTATAAAAATTGGCAAGAGTATATCTGAGTA

Fig. 9.133

AATGTGACATATGAAAACATATGTTTCAAATATGGCCTTAAATATGCAGAAAAATAGGACTTTTCTTCTCTCTCTTC
ATGGGCATGCATGTAAGAAAGATATATATAAAATATACAGTCATGTGCTATATAACATTTTAGTCAACTATGGACCATAC
ATACCATGGTGGCCATAAGATTATAATACTGTATTTTACTGTATCTTTCCTATGTTTAGATATACTTAGATACACAAA
TATTTCCCATTTGTGTTATAATTGCCTACAGTATTTATTACAGTAACAATGTACCATCTAGGTTTGTTTAAATACACTGT
GATGTTTACACAAGGAAATTTCTTAATGATGCATTTCTCAGAATGGATCATTATCAAGTGATGCATAACTGTATGTGTG
TGTGTGTATATATATATATATATATATATATATATATATATATGACACATGCATATGTCAATGTATATATTTAATGATGTTTGCAAT
TCATTTTCATAGAAGTTATAAAAACATTTTGTATTTATCCTTGATTTTATAGAATTCACACACAAAGCTAATAAATTCTAA
GATTACTGAAGATGTTAGTTAACTGTTTTAACCAAGCCACACCTGGTGAACAAATTTGCTATTTTAAAAATAAATTTAG
ATTTTAGTTAATCAAATATTGTTTCAGTTAATGTCACTGTTTTCTAGATAGCATTTTTCATGTATACATTTTACTTAATT
TCTTTCCAGGAAAGTATTAAGCCATTATGTAATGATAAACAGTGTGATTTTTGTGTGATTATATGTACTTTCTTGAATT
ATTACCTCAGGCCTCTGAGGAAACACTACATTCAGTAATGAAGAGGAAGACCCTTTCCGCGGAATGGAACCTTATCTT
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TGACATTTTGAGGTATTGCTTACCAGAATACTATTTTCAGCTCCAGTATAAGGTGATTTTATTTAGCATTAGTACTTAT
AGAGGAGTCAGCCACAGCCACGTCTCCATCATGTAACATTTTTATTTCTACTTTGCAGGGTATAATTTTGTTTTTAT
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GCCAGTGGGCCATACATAGATAGACTCTTGAACATTTGCTAAAACACAAGAACAACAAATTAATGCTACAAATGT
GATGGTCCACATTTCTCTGCCAAGTTAAAGGCATCTCTGGGGAAAATGTCTTTTGATCTTGTAAAGTTAGGAAGGTT
GTGCTCAGAGGAAATTTGGCTAGTAACCACTCAAAGATACTGAATTTAGCTGTTTTACTTGTATTGACCAGTCTAA
GGGACCTGTGTTAGAAAAAGGTATGAGGAAAATACTACTAGATAAATAAAGAAAACAATATTCACCTTGAAGCAAT
GAAAAGAGGAAATATATTTTGTCTGTTGTTGTAATTGAACATATGTGTTTAAAGAAGAATGCACACACATTGATCAGA
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CCCTGATATATTCAATTAGCTTTGGTATCTCAAAAATTTTGGTTGCTAACCATGATTTCTCATTTTGAGTTGGTGTAT
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AAGATCCTGCCACTGCACCCTGGCCTGGGTGTGAGAGAGAGAATGTATTGCTAAAAACAGTTTCCAAGTGCCAAGGAAC
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GGCCAATTGCAAGAATTGAGTATGCATGAATTTTGGTACATGTGGTAGACCCGAAACCAATCCCCTACATATAACCAAGG
AATGACTATGTAAGCAATGTATGCTTACTATTTACAAATTTATCCAATATTATGAAGAAAATAATTTTCTTTTGA
AGATGGAACAAGGGTTTCATTTCAAAGAAACAATAGACAATAATGGCCAATCAACAAGAACCAACTAAGTAGATATT
AAAGGTATTGTTATTACATTTGGAAGCCAGAACAATTTGCGAATTGAATATTGAGGACTAGAAAATTCAGGAGTATAAC
TTTCTGGAATGTGCAAGCTTTGGAGGAGAAGTAGAAGGTAAAGTGCGGGGCCAGGTGCAGTGGCTCACGCCTGTAATCC
CAGCACTTTGGGAGGCTGAGGCGGGCAGATCACAAGGTGAGGAGATCAAGACCATCCTGGCTAACACGGTGAAACCCAG
TCTGTACGAAAAATACAAAAAAAATTAGCCGGGCTTGGCAGTGTGTGCCTGCAGTCCCAGCTACTCAGGAGGCTGAGG
CAAGAGGATGGCGTGAACCCGGGAGGCAGAGGTTGCAGTGAGCCGAGATTACGCCACTGCACCTCAGCCTGGGCGACAG
AGTGAGACTCTGTCTCAAAATAAAAAACAACAAAAACAACAAAAACGAAAGGTGAAGTTCCGGGCTTATATTCTGTA
GCCATTTTAGTTAACCAGGAGCCCCAGATTCCTTTACAGCTGTGAGAAAGAGCTAATGGGAAATTTAGCCTATTTTAAC
TGAGTTGTCTCCAGCCCATTGCCCTCATTTAAAGGTTAGTATGCCTTTAAGTCCCATTTGCCAACCTATACTGAGACCA
TACCTTAGCAATTGCAAATCTTTGATGGTCAAGGCGACTTTTCATGCCCTCCAGATTTATTTTCTTTCTTTCTTAACA
TCATAGCCTTTTGCTCTTTCTTCTGGCATATCAGTATGCCAGATACTTCTTTGTCAAGTGTGAGGCAGTGGGTCTAGA
GACTTGTACCACCCCTTTAGAAAGCCACAGTATGGCCCTGAGAATCATTAGTTTCTGAAAAAGAACTTTGATGTGCAGA

Fig. 9.134

Fig. 9.135

AAAAATATATCACTTCTTGGGTAAACACACTTCCCCCAACCAAAAAAAATTCTAGGTACTTAGAATTCTTTCTAG
AACTTAGAATTTCCATATTATTTTTCATCCCATTCTCTCATTCATTTAGAATCTGCCCTCCAGTTGAAAGGAAGTAGA
TGGAGCCGGGCACCTTTTAGATTATTTCAAATCTGAGGAATTCAACAAAAATAAATGCTATTTAGTTATGTTGTTTGAAG
TCATGTATGCAACATATCTGGATGATGGGTCAACTTTGTTTAATAGATTAGCATCATTTTACACTGAAGTTCGTGGCTCT
GAGCTTCGTTAAAGTGAGACCTTTGTCCACATTCCCAGGAGGGCTTTTTCTCAGGGGGTTCTGAATTTTACCATGACGT
AATTCCTAAACAGTTGTGATTAGTGCTAAATGTTATCATAAACACACACACACACACACACACACCCCTTGCTGTGTGAAA
GCTAGATATAGATATTAAAACTTTAGAATACTTGTTTTTTAAATATGTAATTTATAATATATAAATAATATTTAGAAT
ATAAAGTTTATAATACATAAATACTATACTTCTTTACCATTCACTTTCTAGAACTTTCCAGCTTTTGATCACATATTT
CAAGGGTGGTCTCCCTCTTTACTCCTCTACTAACTCTTCTGTGTCTCAGTTAACATGTGATTTTGGCTTTCTTATTTCT
CCAACATTATTTAATTCAGACTAACTTTATTTCTTGAAGGTATATAACCAGACATACCTCGGAGATATTGTGGGTTTCAGA
TCCAGGCCACTGCAATAAAGCAAATATCACAATAAAGCAAGTCACACATACTTTTGGCTTTCCAGTGCATATAAACT
TATGTTTACACTATATTGTAGTGTGTAATAGCATTGTCTCTAAGAAAACAATGTACATGCCTTAAGTAAGAAATACTTA
ATTGCTAAAAAAAATGCTAACAACCATTAGAGCCTTCAGCAAGTCAGAAACATTTTGGCTGGTGGAGAGTCTTGCCCT
CAATGTTTCATGACTGCTGACTGATCAAGGTGGTGGTGGTGTGAAGATTGGGGTGGCTGTGACAATTTCTTAGAATAACAC
TACAGTGAAAATTGCCACATCCATTGATTCCTCCTTTTCATGAACAATTTCTCTGTAGCATGCAATGGTGTGTTGACAGCA
TTTTACCCATAGTAGAATCTTTTCAAATTTGGAGTCAATCCTCTTAAACGCTGCTGCTGGTTTATCTACTAAGTTTAT
ATAATATTTTAAATCCTTTGTTGCCATTTCAACAATGTTTACAGCATCTTTACCAGAAGTAGATTCTGTCTCAAGAAAC
CATTTTCTTTGCTTCATCCAAGAAACAACCTCATTCTTTCAAGTTTATCATAAGATTGCAGTAATTTTATTACAAC
TTCAGGTTCCATTTCTAATTCTAGTTCTCTTGCAATTTCCAGCACATCTGCAGTTACTTTCTCCACTTGAACCCCTCAG
AGTCATCCATGAGGGTTGGAATCAATTTCTTCCAACTCTTGTTAATGTTGATATTTTGACCTATGCCTATGGATCACA
AATATTCTTAATGGCATCTAGAATGGTGAATCCTGTCCAGAAAGTTTTCGACTTACTTTGCCCTGATCCATCAGAGAAA
TAAGTGTCTATCATAGCGATAGCCCTATGAAATGTATTTCTTAAATAATAAGGCTTAAAGTTGAAATGACTCCTTGGA
ACATGGGCTTCAGAATGAATGTTGTGGTAGCAGGCATGATAACATTAACCTCCTTGTCATCTCCAGCAAAGCTCTTGA
GTGACTAGGTGCATTGTCAATGAACAGTCATATTTTGAAAAGAATCTTTTTTTCTGAGCAGTATGTCTCAACAGTGGGC
TCAAATATTTAGTAAACCATGCTGTAAACAGATACACTGTCATCCAGTCCTTATTTTTTTTATTTATAAAGCACAGGCA
AAGTAGATTTAGCACAAATCTCAAAGGCCCTAGAATTATTGGAATATTAAATGAACATTGTCTTCAACTTAAAGTTACC
ATCTGCATTAGCTCCTACCAGGAGAGTCAGATTTTCTTGAAGCTTTGAAGCCAGGCATTGACTTTTCTCTCTAGCT
ATGAAAGTTGTAGATGGTATCTTCTCCAATAGAAGGCTATTTTGTCTCCACTGAAAATCTGTTGTTTAGTGTAGTGCC
TTCATCATTGATCTTAGTTAGACTTTCTGGATAACTTGTCCACAGTTTATACATCAGCACTTGCTGCTTCACTTCACACT
CCTTTTTTTTTTCCAGCTTTTATTTTAGGTTTCAAGGGTATTTATGTCAGCTTTGTTAAATGGGTAGCTTGTGCACCCTGT
ACTTTTATGTATTGGAGATGGCTTCTTTTCTTTAAACATCATGAACCACCCTCTGCTGGCATCCAATTTTCGTCTGCA
GTCCTTAACTCTCTCAGACTTCATTGAATTAAAGAGAGTTAGGGCCTTGTTTGGGATTAGGCTTTGGCTTAAGGGAATG
TTGCGGCTGGTTTGATCTTCTATCCAGACCACTGAAGTTTTCTCCACATTATCAATAAGGCTGTTTGGCTTTCTTGTC
TTTCTGTGTTTACCAGAGTAGTACTTTTAAATTTCTTCAAAGCTTTTTCTTTGTATTCACAACTTGGCTAACTGGTAC
AAAAGGCTGGCTTTTCAAGCTGTCTTGGCTTTTGACATGTCTTCTCACTAAGCCTTATTATGTCTAGCTTTTGACTTA
ATGCGAGAGACCTGTAACCTTCTTCTTCACTTGAACACTTATAGGCCATTGTAAGGTTATTAATTGGCCTAATTTCAAT
ATTATTGTGTGTCAGGGAATAGGGAGACCTGAGGAAAAGAGAGAAATTGGGGAATGGCCAGTCAGTGAAGCAGTGAGAA
TACATGCAACATTAATCCATTAAAGTTTACTGTCTTACATGGGCATGGTTTGTGGCACCCCAAAAATTAAATAGTAACA
TCCAAGACCACTGATCACAGATCACCTAGCAGATATAATAATAATGAAAAGTTTCAATAATTTCTGAGAATACCAAAA
TATGACATAGAAACACAAAGTGAGTACATGTATTGGAAAAATGGTGCCAATAGACTTGCTTGATGCAGCATTGCCAC
AATATTCAATTGGTAAAAGCACAATATTGTGAAGTGTGATAAAGCAATGTGTTAGTCTGTTTTGCATGGCTATAAAGGA
ATACGTGTGACTGGCTAATTTACAAAGAAAAGAGATTTATTGCTCATTTGTTTAGCAGGGTGTCTGGGCAAGGCACCA
GCATCTGCTTGGCTTTTGTGAGGCTTCAGGAAGCTTTTATTTCATGGCAGAAGGTGAAGGGGGAGCAGCTTTGTGATG
ACAAGAGAGGGAATAAGAGAGGGGGCAGGTGCCATATTCTTTAAACAATTAGATCTCACAATAACTCATGACCACAGGG
AGGGCACCAAGCCATTTATGAGGGCTCTGACCCCATGACCCAAGCATCTCCCACTAGGGCCACCATCAACATTGGGAAT
CAAATCTCAACATGAGATTTGGAGGGTATAAATACCCAAACCATATCAAGCAAAGCACAATAAAACAAGTATGCCTGTA
TTTATACACATATTACAAAACAGAATTAAAAACAATAAAATCATACATGCAAGGGATTCTGGTAAATATACTGATCCCC
TCAATCATGCATTAGTCATATATTAATGTATTCACTTTATATTCAACCAACAAATAGTGAGAGTCTGATATTTGCTGTA
TACTGAGGTATGCACTGGCGACCCAAGGCTCAACAAGACAAAACACAGCCTTTTCTTTTACAGGGGATTGAATCAAGTA
GGGCACGTGCAAGAAAATAGACCACTATAATCTTGTTTGTTCATATAATGTACAAATATGACTGCAGACAAATGGAGA
AATATGAAATCTCACTGGGGAGCAAACCTCACTTTGGCTTGTTGAGGAACTAACACCTAACCTGAGACCAAAATAAGATGG
TTCCAGGAAGAGAGAACAGCAGGTCAAAGTCCAGGGATGAGATTAAAGTAGAGTGTCTGCACTTAGCTAAATAAAAAC
GCTAAAACATGGCTAGACCAGAATGCCAATTGAAGGTACAGTTGATAAACTTACCATTAACTTTTCATGTAACCAATA
TGTGAGGTTTTGTAATATTTTCCAGTATGCTAAGCAGCTATGATATAAACTGAAAAGGCAAACAGTTGGATTATTCC
AAAATTGGAAAGCAGGTGCAATATAAGAAAAATAGGGCAAGGGATTGTAAGGTATTTGAAAGAAAGTGATAGAAATGAT
GAATTGTGGCAACTTAGCTGGTTATGGAATGAGTGAATATAAGAAGATACTGATATAGAAAGTATTCAAGGTCTCTAG
AAATCTCTATGATGTCAAACAGGTGAATTAGAATAACTGAGCAAACAAAAGATAGATCATATTTGGAGGACAGAATGAA
CTTTAAGATGTAAGTGCTAGAGGTAATTTAAGGTATGTTCAAGACATAGAAGGGTGTGAGGGAAGAGGGTGACTGCCTT
GGAAAAGATGTAGCTGAGGAACTGAGAGATCGGGGCATGAGGTGATGCCTCCACATCAACATCAAAGTCACCCAGGACT
GCAGCAGGATTCAAGCTGTCAAGATGACAAAAGGTGGGCAGTTTTTCAGTGAATGAGAACTGACCAAAGGGACTGGCA
GAAGTCAGAGACAAGGAAGTAGAGTTGATTGGCATAGATGTCAAACGAGAAGTTTTTACAAGAAGGTGGAGGAGTAAGT

Fig. 9.136

AAAAGTGAGTAACCAGGAGAGCACCCATTTTCATCTTGGGACCCAGAGAGGGCATTGAGAACCAGAGTACCTTACCCTGA
ATCCACCACAGGGAAGTGACATCTTTAGCGAAGAGTCAAATTTGAGAGAGGTTAGGATCTGGGATGACTTTTTTGGTGGC
AAGGCTGAGGATGAAGTGAAGTCTGTTTGCCATGGGAAAAGGGATCCAGAGAACTGATTGAATAATATGGAAAGAATG
AGAATGGGAGGATAGACAGGGGAGAGAAATCACAGTGCAACATGGGGATAAGAAGTTGAAAGAACACAGGGAGAGGAAG
AAGCTCTGTATCAGGTGGTTGTCTGAGATATGATTATGAAACAGCATTCCAAAATATTTTCAATAATTGGTCCACCTGT
CCTCTGACTGATGGGTGTTAAACCAGGGAACAAGAAGTTGAGCATAGCTAGTAGGTCTTTTGGAAAGACCATAACGTTT
GTCCTAGTCCCAGTAAGTAGGTCTGCAATGGTGTCTTGGGTTCCTAGTACATTTTCAGCATCTATTGGAGAATAGGAGC
TTCTGGTGTTCAGTACTTCTGCATCTTCCAGATTACTCAGCTGTTTCCCAGATTTATTACAGTCTCTGGAGTATGGAG
AAGGATCTGATTTTTTTAAAAAATATGAATTATCATTTTTCAAAGTATTAGTTTATTTTTAAATAATTTTTATATCTAACTC
TAATATTTTGCCAGCAATAATAACATCTTAGGTATTTTGAATTAGCATTGAGTTTGCTTTTGAGAATTTTAGTTCTTA
GAAATAATAATTGAATTAAACACTATTGCTTTCTTATATATTCTTAAATACTATTTCTGGCATTCAAGCATTATTCAG
TTCGTATGCTGCCAATTTTAGCAATAAAATAAAGAATAAAATTGAATTCAATGTGGAAAAATTCTATTTGTAGAAAAC
ATTCGTCTGTAAACGGGGTTACTGTTGTGACCTTAGTGAAACAACAGAAAGGCACGTGCATACGGTTTTGCCCCTTAGCT
TGGATGTAACATATTATTTGCAGACTTTAAGTCATAGCAACCTATGATCATGCTAATTTTTTATTTAAGGCTATTTGATA
GAGTTGTATGATTGAGACCTGGAACCAAGATCCTTGATGTAATACATATGTGCACACAATTTTTTTTATTCTCTTCTCTT
TGGTGACCTTGTCTGGGTCAATTTTTTTTGTGTTTGTGTTTAAATTTACACTGTTTGAAACAGTATCTGTGACTAGAATTTG
GCAACTGTAGAGTATATCATATGTACACAGTGCCTTAGCAAAGGGTTTTAAATTAACTCAGTGGAGGAGACCCAAGCAG
AGGAGCCTTCATTTGTCTTAAGAAGAAATGTCAAGGTTAATTGATTTTCAGAGGATTATCTGCAACAAATAAAGACCAA
GACTCATATCTTCTTATTTTTGGCTTATAAATTCTGGTATGTTCTGTTCATATCCACTGAAAAAACTGAGATGAGAGT
GTTTCTATAAACTCTATATGTCAAAGCATCCTTTGCAGTTGGCAAACCTGAATGAATGACAGTGAAAATGTATACTTA
TGGATTATGATATAGTCATTAAATGATTATTATGATAAGTATATGGTAACATGGAAGTATTTAGAATGTTTTGAGGAG
AAAAGCAAATATACAAAGTATATCCAGGTTAAAGTTAAACACTGCAAAATATATGCCTGCATGTGGCTGAGAACTGGA
AAACAATGTGAAAATAATTTGAATTTTTTTGATTTACAAAATTGTGGTTTTTTTATTTTTCTCACATTGATTTCTATTAAT
GAAAATAAGTGTTTATCCAATCAATAAAATTACATTAAATTTTATTATTTTACAATTTCAATTCATACAAATTCATC
TATTGTGTCTTAGTATATGACAGACACAGTTTGAGGTACTGGGAATATTTTATATGTAGATTTTAAAAATTCATGAACT
TTTGTTTTTTCACAGGCAATCCAATAGGAAGTTTTTGGCAACTATGGTGGGAGGTAGGGATACTTGGGCCTTTCAGAATTC
TCCAGATCACTTTGTATAACCATTCCTCTGTTGGTGACCGACTCCCCAAAAAAGATCACCAGCAGAACCAAGTGAACT
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ATATTTGTAGAGTTTTATAGCCAGGTCTGGGAGCCTGATCTTGCAAGGTGGAAAAATGGCTGGCATTAGGCAGAATTTA
GAGCCTAGCAGTTCTGGATTAGTGGGGTCTCTAAGAGCAGTGAGGTTCTCAAAGTGAACCTTGATAAGTGAACCTGTTAG
TCTTATTGAATAAGCTGGTTCAGGCCATTTACAGCCTTATTTTTCCAAAAGCAAACATTTCTGGAGCAAGTACCTAAGT
TATTTTTGCTTGGTCTCCATGTGCTCAGCATTAACACAGGCATTTAAACTTTGACTCCAAATTGTTTAGCACTGGGACA
GAAAAGTATTTTGGTTTTAGTTTTTACCTTCTTTGTGAGCGACCTAGGTTTGATCAACTCATTGCGCTGACAAGCCGGA
TCCTTTGTCCCTTATTTTTATTTTTTTAGACTTATATTATGTCACCTATATTGCTTGGAAATTTAGGAAACACAGTCCT
TGGGATAGCTGGGGGGATGAGAAGTCAAGAGAGGTAAGGTCTTAGGGTAGACAACCTGGGTAGATACTGGGCTACTCAGT
ATCCAATCCTCTTCTGTTTTTGGGGGAGTTTTCAGAATAGGTGAGAGAAAGGACCTAGCCTTCCACTAGGGGAGTAGGAAT
ATGCATTCCCTCTGCATGCCCTCTGGCAGACAGAGGGAGGGTCTATAACCTCAATGTAGTCAATCTGATGTTCCCTAGC
CAGGACTTTGAATCTTGAGGGGATGGCTCAGACCATCACATATGCCACTATCGAGTGGAAGGGCCAGTGGCATCTGTTG
TAGCTGCTGACAGTGTGGCGAAGAGCAGTGTGTGCCAGCAGGTGAGGAATAGGGTCCCAGGCAGACTGATCCAGCCTC
CCAGCTGCTTATTATCTTCACTGGATTCTGCTCAAGTCTGCTTCTTCAAGCTCCCTGTTTATTCTGAGTTCTATCCTAT
ATTCTTTTGATGGATTCTTTTTCTTGCTTATGTCAATCCATCATTTCTATTGTTTGCAAATCACAGCCCTTGATCACTC
TAGAACTAAAGTACAGAAAGTGCCCTGACTTTGCTAGAAGTTTAGGTGTAAAATTAGTAGTAAAGTTAGTAAGGAAAA
ACCAAAGGGGAAGATTGAATTTTTATTTTAAATGCATTATTTTTTTTACATAAAAAGATAACAAAAGGAGGGTAGTGAAGAG
AAGAGTCTCTTGAGATGTCCGTGGCAAAGTTTTTAAAGCTCTCAATAAGGAGACAAGGACTGGGCTTCATTGCAATTAT
TCCCTTTTGACTGTTGAAGCTGGTAGGCTTACAATGGACTCCTAATGAGAGTGGTTTTGCACTACAATTAATTAAGTGCA
TGACATTATAAACTGCATGTCTCTTCCCTCTGGAGAATATTCTTCCCTATATAAGTGCTATTTTTGTTTTAATTTCTCTCT
CTGTCTCTTCTCTCTGTTTCCCTCTCTTTGTCTCTCTCTCTTTTTTTATTTATTTATTTATTTTATTTATTTATTT
AAGTTTTAGGGTACATGTGTACATTGTGCAGGTTAGTTACGTATGTATACATGTGCCATGCTGGTGCGCTGCACCCACT
AACTCGTCATCTAGCATTAGGTATATCTCCCAAAGCTATCCCTCCCCCTCCCCCACCAACAGTCCCAGAGTG
TGATGTTCCCTTCTCTGTGTCATGTGATCTCATTGTTCAATTCCCACCTATAAGTGAGAATATGCGGTGTTTGGTTTTT
TTGTTCTTGCGATAGTTTACTGAGAATGATGATTTCCAATTTTCATCCATGTCCCTACAAAGGACATGACCTCATCATTT
TTTATGGCCACATAGTATTCCATGGTGTATATGTGCCACATTTTCTTAATCCAGTCTATCATTGTTGGACATTTGGGTT
GGTTCCAAGTCTTTGCTATTGTGAATAGTGCCACAATAAACATACGTGTGCATGTGTCTTTATAGCAGCATGATTTATA
GTCCTTTGGGTATATAACCCAGTAATGGGATGGCTGGGTCAAATGGTATTTCTAGTTCTAGATCCCTGAGGAATCGCCAC
ACTGACTTCCACAATGGTTGAAGTAGTTTACAGTCCCACCAACAGTGTAAGTGTTCCTATTTCTCCACATCCTCTCC
AGCACCTGTTGCTTCCCTGACTTTTTTAATGATCACCATTCTAACTGGTGTGAGATGGTATCTCATTGTGGTTTTGATTTG
CATTTCTCTGATGGCCAGTGATGGTGAGCATTTTTTTCATGTGTTTTTTGGCTGCACAAATGTCTTCTTTTGAGAAGTGT
CTGTTTCATGTCCTTACCACTTTTTGTATGGGGTGTGTTGTTTTTTTTCTTGTAATTTGTTTAAAGTTCAATTGTAGATTCT
GGATATTAGCCCTTTGTCAGATGAGTAGGTTGCGAAAATTTCTCCCATTTTGTAGGTTGCCCTGTTCACTCTGATGATA
GTTTCTTTTGTGTACAGAAGCTCTTTAGTTTAAATTAGATCCCATTTGTCAATTTTGGCTTTTGTGTCATTGCTTTTG
GTGTTTTAGACATGAAGTCCTTGCCCATGCCTGTGTCTGTAATGGTATTGCCTAGGTTTTCTTCTAGGGTTTTTATGGT

Fig. 9.137

TTTAGGTCTAACGTTTAAAGTCTTTAATCAAAAGTTGTCCCTGAAGGAAAATTAACCTCTGCTGTTAGTCCACTGAGAAAG
CTTCTCTTTTAAATACTGGAACATATCTTTGTCTAACTCTCCCTCACACCTAGTGCTAAATTTATCAGCGTTGTTCCCTGA
TACAAACATTTTGAAGACAATCGTGAGGAAAAGATGAAAAAGTTGCATTTAACTTTTGTACAGGTGTCCTTGTTCT
CCAGAAGTTACTGTCTTTGTTGTAGTGTGCATGCAGTCACAGGAGTGTGCATACATGTGTGAAGTACTTGAGAAGTGTA
TTTGTCCGTTCTCACACTGCTATAAAGAACTGCCCTAGACGGGGTAATTTACAAAAGAAAGAGGTTTAAATTGACTCACA
GTTTCAGCATGGCTGGGGAGGCCCTCAGGAACTTACAATCATGGCAGAAGGGGAAGCAAACCTTCTTCACGTGGTGGCA
GCAAGGAGAACTGCAGAGCAAAGTGGGGAAAAGCCTCTATAAAACCATCAGCTCTCATGAGAACTCACTATCACGAGAA
CAGTATGGAGGTAACCTGTCCCCATGGTTCAATTACCTTCTACCAGGTCTCTCCTGTGACACATGGGGATTATGAGAACT
ACAATTCAAAAGATGAGATATGGGTGGGGACACAGCCAAACCATGTCAAGAAGTATTTTCAGAGACAAGTCATTCCCCAA
ACCTTATCATCAGGGTCAAGACCTGGCAGGTGAGACATTAGCAGGTGAACACAACATTCTCAGAAACCTTGAGAGTAC
TCCTTTCCCTCCAGGCATCTTCTTTCAACCTTAATATATTCTATTCAATTATATTCACAGCTTTTACCTCAATTATAA
TTCCATTCTTCCAAATTTGTTTTCTTCTCCCATCTTTCCTCCTAATAAAATATCTATTCTTCAGATAGTCTTCCTTTGT
AAAATAGAAAGGATCTACAAAAGCCATTCTTTGTGAGATGTGAATCATTTCAAATAAATTAATGCCTTGTTCTAATGTG
AATGGTGGACTTGTGAAGAGAAGTGGGTAGATATTTGAATTTCTGAATGCTTTAGAATATTAGTTGCACATGCAGTAAT
ATTTCTGTAGCTTAGAAGAAATTGGTTGGCTTAACAGAAAATGCAATTAATAATAGTTTACAAATAGGTTCTGGCATCA
ATAATAGAAATATTAAAGAACTACATATTATATCAAGATGCTTCTCTTTTTTTTCGGGGGGGGGGTGCCTTTTCTTCT
TGATGAACACAGTACTCTTCTTCTTGTCTTTTATCTTCCATCTCAAATGTCAATAATAAATTTGAAGAATGGAGAA
TAATACATTATCACGTGCCAGACCTTATGCTAAACACTGGATATGTTATATTGTGTTTCATACTCAGGATGATGCTACG
GTAGGTACCATTATTATCTTATTTTACTTATAAGAAAACAGAAGCCTGGAGAAGTTAAACAATTTCTCAGTATCAGAA
AGAACCAAGATCAAATATCTAGGTTAAGGTATTTTATCTTAACCTAGATATGCCAGAATATCAAATCTAGGTTTCACTA
TTATTTTGTATTCTATCCAAATTTCTAAAACCTGCTAATGATGGAGGGCTGTTATATGGTTCTAGCTTTATATATTTTTT
TATTTCAACTTTTATTTTAGATATATGGGTATTTGTACAGATTTGTTACATGGGATTATTGCATGATGCTTAGGTATGG
TATATCCCATTAACCTGATAGTGAGCATAGTACAAGATAGGTAATTTTTTAATGCATCCCACCCTCCCTCTACCCTCTA
GTAGTCCATGGTGTCTATTGTTCCCATATTTATACATATGTCCACGTGTGCTGAATGCTTAGCTCTCACTTATAAGTGA
GAATGTGCAATATTTGGTFTTCCATTCTGTGTTAATTTGCTTAAGAATATGGCACCCAGTGGGCCGGGCGCAGTGGCT
CACACCTGTAATCCCAGCACTTTGGGAGGTGAGACAGGTGGATCACCTGAGGTGAGGAGTTCTAAACCAGCCTGACCA
ACATGGAGAAACCCCGTCTCTACTAAACTACAAAATTAGCCAGGCGTGGTGGCGCATGCCTGTAATCCCAGCTACTCG
GGAGGCTAAAGTAGGAGAATTGCTTGAACCCAGGAGGTGGAGGTTGTGGTGAGCTGAGATCGGGTCATTGCACTCCAGC
CTGGGCAATAAGAGTGAACTCACTCTCAAAAAAAAAAAAAAAAAAAAAARGCATATGGCTCCCAGCTCCATTTCAT
GTTGCTGCAAAAGACATGATTTTATTCTGTTTTGGGGTTGCATAGTATTTCTATAGTATATATGTACCACATTTTCTTTA
TGCAATCTACTATTGATGGGCACCTGGGTTGATTCCACATCTTTGCTATTGTGAATAGTGCAGTGATGAGCATATGAGT
GCATGTGTCTTTTGTAGTAGAATTATTTATTTTGTGGAAGTATATACCTGGTAATGGGATTGCTGGGTCAAATGGTAAT
TCTGTTTTAAGTTCTTTGAGAAATCTCCAGACTGCTTTCCAAAATAACTGGACTAATTTACATTCCCACCAATGGTGTA
TAAGCATTCCCTTTTTCAGCCTCGCCAGTATCTGTCAATTTTGTACTTTTTTATAATAGCCATCCTGACTGGTGTGAGAT
GGTATCTTATTGTGGTTTTGATTTGCATTTCTCTGATGATTAGTGACGTAAGCATTTTTCATATATTTCTTGCCACTT
GTATGTCTTCTGTTTCATGAAGCCCTTTGCCCCACTTTTTAATGGGGTTATGTGTTTTTTGCTTATTGATTTGTTTAAAGTT
CCCTGTAGATAGTGAAAATTAGGCCATTGTCATATGCATAGTTTGCAAATATATTCTTCCATTTTGCAGGTCTTCTGTT
TACTCTGTAGTTTCTTTGCTGTGCAGAAGCTATCTAGTTGGCCATCTTGGCATCTAATCTTTAATTTTCTTTCTAAAA
TATTTGAATGTTTTGCAATTTCTGACTGAAGTTACTTCCCTCTTTCTGAAGAAGGCCCTGCTGACATCAATAATTATCT
GAGAGTGACATAAGCTGACTCCGATTATGCCAAAGTAACCCTTACGTGGTATGAAAAGAAAATGAAGGTGACTATGATT
CCTGGGGCATTGAGATTCTGAGAAAACCTCCAGGTGAGCTCGCCATAAAAATTTCCCCACACCTGTAGTTTAAATTTACCA
ATTAGGAAATTATATATTTGGGGCTAAATGGTTTCTGATTTGTTGGCCTGAAAGAAATAATGTACTTTTAAACAATA
AAAGTTCTCTGCTAGGAATATCTTAAATACAGTGAAAATCTGCCTTGACAGTGGACAGTAAGTTAGATTTTCAATTGTTGT
AAATTGGCACCATTAACTAAAACCTTGTCCATTCAAATAATATGTATTGTACATAACACTAAGCTCTGTTATTTAA
CTGCCACAGATTATTGCTTTAACAGTTTGAAAATGATTTTGTAGAAAATTGACTTTGTACAATGCTATATTAGAGTCTGT
TAATTGATGAGAAAACCAAACCTCAGATGACATTTGCAAAAATAAAATTGGCATCAGTTGAAACATCTGTACCATTTCTC
TTATTTAAAAGACATTTTTCATCTGGAAAATGACTACTTCGATTAGCCTCAGATCACAACCACAAAATAAAGTTGTCTA
TGGGTGGCTTCTTATTTGCATTTAAATTTATTTATGTTGCAGCATTTTGTATGCACAAGCAAATGGAAGATTGTTTTGAAT
GACATAGGCCTATACACCTGAGAGAGAAATATGTCATCTACAAAATGTAAAAATATAGGTTTTTGTAAATGCCTCTGAA
AGGATTATAAGAATCTACATTTTCCCTCCCTCCTGGAGTTTGTCTTTCTATTAAGGCTCTAGACGAAGAAAAGCTCTA
ATATACTTTCTTGGAGGGGTCATGAGACAATGAAGCATAATAGGATTTGAAGTATATTGGATATAAAATTCATGTGAT
ATCCTCAAGTAGAGAGATGTTATTTTACTGTTCTTTAATCTTTACTGTTGGCGATAATTGCCTTTGTAACTAACTTTA
TGTTGTAACTAGTAGCGAACCAGAGATTGAGTTGAGAGGTGGGGTAACAGATGAGATCCTCACTTGTGAGGGCTCCA
AGGGCCTGTGAGGCAATGGAAAGGGCTGACCTTGCTACCTGGATGGGAACAATGCTGTTATCATTTTCATGTGCTCCTG
TTTAGAAGATGAGTTAACATCCATAAAAACAAGTTAGACTAAGGGTTGTCACTACAGCCTGTGGGCCAAATGCTGCC
TGCCACTTGTTTTATAAATGAAGGTTTCTTGGAGCTTGCACTACTACTGCAGAGTAGAGTAGTGAGATGGAGACCATT
CGGCCTGCAGAGTCAACAATGTTTTAATATCTGGTGCTTTACAGAAAACTTTGCCAAGCCTTGATTTAGACTTAGTAAC
TTTAGTAAGAAGTTACTGACTGCACAAATTACCAGGATATTTAAGCTTGTAAGTATTAAAATACCATAACTATCTGCT
TCTTGTGGCAGGGAGGGGGCCAAGGAGAGGTGAAGTGGTTATAATGCTCGTGCAATTGGTTCCTTCATTCACCAGAGTGA
TTGTGCGCCCTCTCTGAAACAGGGATAATGCCAGGTACTCAAGATACAGCAGTGAGCAAAAAGAGGCATAGTGCCCTCC
TTAATGGAGATATGCATTGGAGTCACATAGCATAATAATGTTGACAAATATCCCTGCTCTCATGGATCAAGAGGTTAGA

Fig. 9.138

GTTTTAGGGTAACCATGGAAGATCTCACTGAGAATGTAACCTTTGAATAAAGGATTAGAGGACTTGAGGAAGCCAACCAT
GCTGCTAGAGTTTCTCTGCAGAGGAAAGGGGAGTGATGAGATATAGGTGAAGGACAATGTGAGCCAAGATTGTTTTTTT
TAAGTTGAGGAATTAACAGCATGGTTGTATGCTGAGGGGGACAATTCAGTGGAGAGGGGAAAAATTGATGTTGCAGGAAA
TTGCTGGGATAATGTCTTTGAGCAGATAAAAAGGGATTGATCCAGTGTACAACAGAAGGGACTAGTCTTGGTTTTGGAT
CAAGGACAGTTTATCCATAGTAAAATAAGAGAACGTAGAGTATATGGGCACAGATGCAGGTCATTGGGTAGTCGTGGTG
GTTGAAAATTGTAGAAGTTCTTTTTTTTAAACTTTAATTTTTCTTAGTATGTGCAAAGGCCTTGAAGCAAAATAGAGCA
TGGTATAATCAAAGAATTGAGAGAAGAAAAACATAATCTAACTGTAAATTAAGCAATACCATGGAGCCAGATGAGGCTG
ACGAGGTACACAGGATAACAGACGTCAGCAAGACTAAGAATATGGTAATGCTAAACCAATTAAGAATGGGGTTAAATCAA
ATTTGCCCTTTTTTAAAAAAGATGACTGTGGCTACTGATAGAGATTGGATAGGAGAGTGCAACAAGAGTTTCTGTGGAGA
GACCAATCAGGAAGCTACTGCAGTAATTCAGGTGGGAAATGAAGGCWTCATGGTCTAGAGTAGAGGGAGTGGGGAGAAG
AGAAATGATCAGATTTTGAGAGATATTTAGAGGCAGAGTAGATAGACTTACCTCACTTATGTTTGGGTATAGAAGACAT
TGCCTATGTAAAGACTGAGTAGAGAAGAGAGCCTAGGAGTAGATGTGGACACTCAGGAGAAGGATTGAGATCCTGGTTT
AGAACACTGAGTGAAGAAGCTGGGTATTGTTAAAGAAGAGAGCCAACTTCACAACCTGCCTAGCAGTGAAGAAGCAGAT
CAACAAAAATCATATGGTGCGGAAGGGATACTTGCCATTAGATTAGTCCAGCTGAAGCCAAACGCTGGTTGCTTCTGAT
TTTTTTAAATCCTATAAATATATTGGTTAATTTTATTTAAAAAATTGAACTTTTCTATTCTCATAATGAGACCAAAGTA
AAGGTCAAAATCTTAAGAAGAGAATGGGTCAGAACATAGCAGGCAGATTGTTGTTTGACATGATGTCAATGGACACTTC
AGTTGTCATGTTCTAGGCCCATTAATTGGCCTTTCCTCTTTTGAAGCTTCTCTCATGGAACCAAGTGCATTTTATATC
TGCATCTGTTCCAAATATGGGTATAACACAGACTCTGTGAGTAGCCTGCAATGTGTTTCCCTTCCACATATTGCTAGAT
AACTCTGCTTATAAAAATCTGCTTGCAAAAATCATAGAACTTGCAAGATCCTGTCAAAAGAATTACTAFTAGATGAAGTT
AAGAAATGACTGTAGAAGTAAAGAACAAGAGCTGGTCTTAGGAGTTCCATCCCTTCATCAGTCTAGGAGGGTTGGGAT
CTTAGAGGCATTTTTGAGCTGAAAACCTCAAGGCAGGAAACAAGCCATCCCCTCTGAATAGGGCTGTGTTTCTCACCTG
AACAGTGAGAAGAGGGTGTATATTTTCTAATAAAATCCACATTAAGTTAGACAAAACTCTGTATAACTATGGAATA
GACTTTACTAATGGAAGAGTAATGTGAAAAGCTAAATAGGAAAATTTTAGGCACGTATTTTCACAATAGATCTTGCTG
TACTTGGGGGAGGCTTGTCTTCTTGATTACTAAGGTTAACTTAGTGCTTCAGAAGCCAGCCCGTTTTTATGTTCTGCTG
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AGGGAGTGTGAGTGTTTTGTGTCATGGGCGAGCACATACAGATGTTAGGTGACAGTGGTTATTCAGTCATGCTTTCAGAC
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AAAGGGAAATCAAATCTGACTTCCACCCCTCTGAGATTTTTGTTGAGGACACAGAATGAGAGCATAACCTTCAGGTCCA
GAGTGGAATATAGAGAGATCTTCCACTGACAATATGCTATCAACAAAAGCAAGGTAGAGAGGAAACGTTGAGCCTCAGC
AACCTGTAAGTCAAGACCTCTCCATCATCCCTAGATGCTTTTGCTCTGTCAATTGGGATTGTGTCCCTTTCTATAACAC
TGTGACTTCTTTGGATCTCCACAGCCTGACTGTGTTACAGTTCTCCTCTCCAGTCTTTTAGTTTCTCTTACAGAGCTGG
TGCATTTCTGTGGGGACAGAGCAGCTGGAACCTGGGCCCCGAAGCTGGTTCAGATATGGAATTCCTTTAAGAGCAAAAA
TAACAATTAATAATAATAACAAGCCCAATAACAAGTGAATAGTTAAAAGGAGAAAAGATATCACTATGAATTCAAATT
TTAAGTTAACATATATCCCACTACCACAAAATATTATAAATCATAACATTTTATTAACCTGCCCAACCCAATTTTATAA
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GAGCCTCGGCAGGTACTTATGCAAGCAGAGAGCCCTGCAGCTTTGGCTTTTATCATCTTCACCATAAACCCATCTCTGA
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TTACAAGTAAGTTTACATTCTTATTCGGGTGAACCTTATCTTTTGTGGACCACAGTTCAACTCACTACAGAAGTTACAA
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CTATAATTTATCTTCTCTTTTCAATTCACATATAAATATTTTTTTTAAATTTTTTAAATTTTTTATTTTAGATTTCAGGTGG
TACATGTGCAGGTTTGTACATGGATATATTGCATGATGCTGAGGTTTCGAGTTTCTATTGATCTTGTCACTAAGATAAT
GAACATAGTACCCAATTGGAAGTTTTTCAGCCCTTGCCCTCTCTCCCTCCCTCTTTTTGAGTTCTTGTGTCACTG
TTCCCATCGTTATGTATGTGTATGCAAATTTTAGCTTCCACTTATAAGTGAAAACATGCAACATTTGGTTTGCTGTT

Fig. 9.139

Fig. 9.140

AGGAGGGAGGGAGGGTTGTAAAGCAAGGATGTGGCTCTGTGTAACCGGGATAAAAGCACCCCTGCCTTAGGGAGATGTTG
TGAGAAATACCCAGCACCCCTCCTGCTACAAGAGGCAATCCTTATAGCTGGCTCTTGCCATGGTTAAATGATTGGGAC
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TGGGCAGAGGAATGGATTTGGATGGGGCAGCATGTTCTGTGAGTTGGCAGGCACCCACTGAACAGAATAACTGATTTG
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AATAACTGCCAAGATGTTTGTGTTAGTAATCCTGAAATAGAAGATTGGCTTCTATGCTAAGATAATTATTGGGAGAATCT
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TGGTTTCTGTGAGTCAAGAAATTGAACAAGGTAGGGTAGAGATAGCTGGTCTCTTGTATGTAATGTCTAAGGCCTTAG
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GAGACTTCAGTTCATCTCCATGAGTTCTCTCCTTGAGGGTTAGTTTGAACCTTCTCATAGCATAATAGCTGATATCCAA
GGGCAAGCATCCCCAGAGAGAGTAAGAACCTGTTAGAATCTGTGTCTTTTTTATGCTTTGGCTTGGAACCCACAGAGCA
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GACCACCATACTTTAGTTTTCTAAACACTGGAGTGCAATCTCTAAATTGCAATGTGTTTGGGGAAAAAAATTCATCCC
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ACTCTTTATAATCCCAAAGAAATATTCCTTAGTTCTTTTTTCTTCTATTTACTATCAATATCTATCACAAAAGCTGATA
TTACATGAAAATATTAATATAAATGTTACAACATTATCTTTCTTACGCAGGTTTGTGTTGTGAGGATTAAATAAATTAATA
TTTATGAAGAATTTGAATCATGCCTAGCATGTAGTATGGACTCAAATATTGCTAATAATATTGACAATTTTTTTTTGAA

Fig. 9.141

ATTGCTCTTCTTCTGTAACACTCAATTTATGCCATTGGCATAAATGTTTTGAGCTAGACAAGATATTGCTCTATTTCAGA
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CGATTTCTTGACCTCGTGATCCACCCGCCTTGGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCGCGCCAGCC
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TACTTTGCTTCAGTTTGAGGATGAAATGAGAGAATATGTGTAAGACATCTGGCACACATAGTGAGCATTCAATAAATGT
TAACTATAAGTAGGTGACCAGTTGGGGCCCAAGGAGATTGTAACCTTGCTTAAGGTCATTAAGGTAGTTTCGTGGCTAAGT
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ATAGGCGTGCACCACCATGCTCAGCTAATTTTTGTATTTTTTAGTAGAGACAGGGTGTCACTATGTTGGTCAGGCTGTTT
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CCTTAAGAGCCAACCTAGCATATCCTTGTAATCAGTTGCCTAAACAAGCTTCCTCTTAAATATGCCTTTGAAAATAT
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TTTCAGTTTTCTCTTGATGCTCAGGATGAACTAAACATCTAGACAAGGGAGCATCTGGGAAGTCTGACTAACTGGACTA
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AGCAAATGGCTTCTATAGTTGAATAATCTTTCAATAAAGTTAGAAAAATGGGTGTCCTTCTTCAATGCATAGCATGTTG
GGAACCCAGGATACTTTAGAAAGTCTCTTGGTCCAGCACTCCGACTTGTGAAAGACAAATATTGTCATACAATGCTTTG
GCTGATCTCTTATTTCTTGTCAACCTAAGGTACAGTGGGTATTAACTCTTTTAGCTTCTTGAAGACAACTAATTCCC
GACCTCCCTGTAATAATTGCTGAACATGGGAAGTTTTGGGGTCTGAGTTTTAATTCCAAGTAATTCATTATGTCTGTCA
TTGTCTGTGGTAGGTGCTTGCCACCACTTTGCCAGTATCATTATCTGTGACTTGAGGCAAATGGTAAGGACAGCTTG
TCTGAGTCTCAGCACTTGGGTGGCAGATGTGACAGAGATAGATAAATTGGCCAGAGTGAGTATTTTAAACCAGAATACT

Fig. 9.142

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GCCCCAAGCCCAGTATGGGACCATCTACAGTATGGAAGGCAGCTCAGACATAGGTCAGTTTGTTCAGGAGACCACACA
TTATGATCTGTTGAGCTGTTTCATCATTTTCACTTGAAAATGAGGGGTGGCTGTTTTTAGGCTGTCAGCCTCAGATTTGA
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AGAGAGACAGGAAGATGATGTTGCCATGTTTTAGTTTAATATAAATTGTTAATATCTTTGTCTCAGTCACTGTATAGAA
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AAGAAGATTAAGATGGAGGGAAGGAGGAAGAGCAGGAAGAGGGGAAAGTAGGTATAGAGGAAGAAGAGAAAAGGAGATAG
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CTCTCAACTAGATACTTTCCATGAGACTCAAAATACATTCAAGTGTCTTCTGCATTAAAAACAAACAAACAAACAACT
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CCTATCACAGCCGATCCTATTCTAGTCTGTCTCTTGTTCCTATTTTACCAATAAAATCATTTTTTAGTAAGAAGACAAT
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CACTTTCTTACCATATTCTCTCACTGTGTCCAGGACACTAGCTCAGACTTTTTCAAATGCTGATTAAAGGGCTTCCATG
CTGACAGAGGGCAAACCTCACAGAGCAAGCTTTTCTTAAGCTTTTGCTTATATCACATTTGCTAGCACTCTCTTGGCCA
ATGCCCAGAGTCACATGGGAAGAGCCTACTGAAGGCAGAAATACAGAGAGAGAAGACTTTTGGCATAACATCTTGAGGG
CTTCCTTTAACCCTACTCCCCAGTAAACTAGTGAAAATATGATCAAACAAATAAAACTCCAAGCACTTAAATCCTCT
GGAAATGATCCTAAGGACAAATAGCAAATGAAGAAACATCTATTCAAGAACATTTATGAAAATTCAATAAGAAAGGCAA
GCCTGTGGTATTTAAACCAAGACTGCTCCCTCTCACCCTTCCAAGTTCAGGGAGATGGAGCTTCCATTCCAGGCTGG
TGCAATCAAGAACACAGAGCTCTCTCCCTCCAGCTGGAAGGTTTCTTCTGGAAGGAACAGAACTTCAGTGTTCCTCA
TCCTGGCCATAGTTACCCATTGCTAAGGCTAAGCTCTGGTGAATACAGTAGAGAGGTAGGGGCTTCCTCCCTGCCAAA
TCCCCCATCATTTGAATGGAGGGGATACCTTAGGCACTGCATGCTAAGAATACAGAGGCCTCATCATCCTTGCTGGCCT
CCTGAGGTGGGGGTTCCACACCAGGAGAGATAAATATAGAAGATATTAGAGTGCTGCCACCTCCCAACTAAGCTAAGCT
CCTAGAGTGGGAGTTTCATGCAGTCATGCAGGAAGAACCTCTCCATTTTCTCCACCTCCATCTTGAGAAACATGGCTTA
GAGATTTTGCTTGGTGGTGGCGGGGCAATAATCCATCATAAATAAATATTCTTAATCTTCTCCCAAAGGGCATGGCT
TCATTTGCAACAAAGTATGGAAAAGTTCAAGCCTTAGAGTGCGCTCAAGAACAGTGGAAGCTATGGTGAAAGGCAATTG
GGAGGAGAGTCAAGATACAGGCTAAATTGCAGACTAGTTTGAGGAGAGAAACCAGAGAAATAACACAGCTGGGAGGAGC
CCTGACTTCGAACATTTCTTCAAAGGAGACACAATTCGATTGTATTAGTGTGTTGAACAATATAAGGTTGTAAAGCAC

Fig. 9.143

TGTTGAAAATAACACAGCAATTGTTACCAATTAGTGGAGTTGAACAGCTGAATGTGGTTAAGGAAAGAGTGAAGGACA
ATCACACGAACATCACTCTCATGCCGGCTGGGCTGTCTGTGGACATACCCAAAGATGTACCTCCCTGATAGCAAACCTCA
GTGACTTAACTGAGGGAAATTAACCTCACCTGACTGTGAAGGAAATATACTTATTAATAATCCAGCCCCCTCACTAAAC
AACCAAGAATAATAGCAAGCCCTGGGTGGTGGTAGGGGGAGTAAGAAGAGTTGCTACAGTATATTATCTGCAATATCC
ACTTTCCAACCAAGATCACAAAGCATGGAAAGAAACAAAAAATATAACCCATACACAGGATAAAAGAACCAGATGGCA
GAAACTATGGGTGAGAGGGACCCAGATGTTGGATTTAATAGAGAGATGGAGAGGAGAATGAATGCAGCCAGTGTGCAAT
GTGTCATGTCCTCTAAATGTTCAAGTCCCTCAAGTCATAGATCCACCTATTTTTTCTCAAGTTAATATCTCTACACCTA
GTTTTAATTCCCTGTCATGTATACAGGACTCACACATTTTATCTTCAACTTAGGCCTTCCCTCTGAGCTCCGGACCTGT
GTATCATTCAGCCTTTTTGAAATCTCTATCCATATGTCTCAAAGGCACTTTTGATTAAATACTACCAAAAATTAACCTCC
CTCTATCTCCCAGTGTCTCTATTTTCAGTGAATGATACCATCAAACATCTACTTTCAAATCAAACTCTGAGATAAAT
CTCTAACAGCTCCCTGCTTATCACCCACCCCATCATATTGAAAACCAAACCCATTTAAATACCTTTCGTTTGTGTCTCT
CCTCTCTTACCTCCACCACCAGTACTGTAGTGTAACTCTGCCATCATCTCTCACATGGATTACTGCACAAGTCTCCTAT
CTGGTGCCCAAATTAGGAGCACCTCCTAACTGGTATCCAATCTAATCTCTACACTGTAGCCAGAAAAAATCCTTTGAA
ATTCAAATCTAATCATGGTATGTGCCTCCCCCTTCTTCTATGGAGACCATAGGAAAGAAACCGAAACTCCTGTGGTGCT
TGAGTTGATGCTAGTAACCTCTAAACCTCATTTTCATGCCTGGATCTGTCACTTCTTGAAGCTCCATGATAAAGCACAGC
CTCCTCACCTCCAAGTGCAGGGGGTCTCTAGTTGTCTCTTCCAGGTGCCACCAGAGGGACTTCTCTCATAAAAGGCCT
GTGCACCTATGCTCTTTCTCTCTCTGTTATGGACTTCTTCCAGAGAGCTGGTCCCCATTCTTGTATGGGACTGTC
CACTTGTATGAGTAATAAAAATCCTTCTCTGATCATACCAAAGAATTGGTGTGATGATACCTCTTTTAAATCTTCCGTCGCTT
TTGAGAGGGAACCTTCCCATGTGCACAGTGAAGGTGTGATGATACCTCTTTTCTACCATCTTTAAATCTTCCGTCGCTT
CCATTCATTTTCTCTTAGGATAAAGTTCACAATCCTTAGGTGGCCTACAAGGTCTCTACAGTTAGGTACCCCCGTC
TGCTTTCCTTCTCTTCTCACATTAGTTATGCTCCCCACTGCCTGAAAACATGCAGACCTGCTGCATCCTCTACCTGAA
ATACTCTTCTCCTCACCCCTGCCAGACTTTACCTAATTTGCCCTTCGTTTGTCTTTGTAACACCGCTCAGAAATCTCTT
CCTCAGAAAGGCCTTCTCTGACACCTCAGCACAGTCTCCTCACGAGGTGAGATGTATTTTTTCTTCAAGGCACTGACT
TTGGTTTAAATTTGTATTTATTCATGTAATTATTTGGAGTGCATTTATCTTCTCACTAGATATAATCACCATTAGCT
CAAGGGCTTTATATGTTTTGCTTGCCATTGTGTCTCAGTGCCTAGGCCATTCACTGACCCACAGTAGGTGCTCTGATA
AATATTTGTTGAGTGTAGAATAAATGAAGACCCTTAGAAGCATAAAATATATAATCTCAAAGCAGAAGATACTTTAGTT
AGAATAAGTTAGGTTCAACTATATTAACACTCCTTACTACACACACATAGCTACACTCATCAATTTTGGTGGCTTAC
ACCAAAAAGGCTTATTTCTTGTGCACACTGCATTCTTGTGCTTTAACAGGATACTCCGCCCTACCTCATCTTTATTACA
GACCTTGGCTGACGGAGGCTACCACTTGGAAATATTGTCTATTTTCAAAGTCCAGCAGAAGAGAAGTTAAGAAGGGTTTC
ACTCTGGCAATTAATAATTTCCAGCTAGGAAATGACATACTTTATCTTCATATACAATCCATTGGCCATAACTAGTCTCA
TGGTCCCTGCCCATTTGTAAAGGGGAAATAGGTAATCATCTCATGCATTGAGGAGGAAAGAAGATTAGATACATGTGAC
CAGTAGAAGTCTCCTTACTGACATTTTAATGTAGAGCTGTATGTGATTAAAGTTAAATTTGAATCTGAAGGGCTGACAG
AAAAATTCAGAGGATATGATATTTAATAATTGAACGAGGCTGAAAGATGAAATTTGGACAGATTGAATGGGTAGGCCT
TGCTAGCAGGAAAGACAAAATAAACATAATCAAGAAGTATGTATGAACGTGTGCCAGGGCAGTGAGGAGTGATCAGCC
TCCTTGAAGTGGAGGATGTGTGTCAAGGAACCAGAGGAGATGATTAGAGTGGCAGACCTGATTATTAGCATATCCATGC
TTCTGTCTGAAGCCTTTTTTGGCCTATAATGTAGAAATATCCTTGGAGTAAGCAGTTTCAGCATATGGTTTTAGACATATA
GGTCATCTTGGCTGGATTGCATATTCTTCATTTTTTAAATTTCCAGGCTGTATTGGTTGATATGCAGTCCAGAAAGAGCT
GGATCTGACTTAAGAAAATAAAAACCTGCTTTAATCCCTCTTTTCAGCCCATTTGTCTATCCATTAGCAGGTAAACACCAG
TGTCCCTGCAAGTGTGTGAACCTGAGCTTCTTAATTTACCTTTAATTGGCACTTCTAGTGACTTAACTGAGTGAAGAG
ACCCTTAGCAATGATACAAAGTGAAGGATGGTATAGAATAATAGGTTTTCTGAAAACCTGAATGCAATTGAAGGGTGCT
ATTTAATAAATCTGTCCCTAGAAGCTAATAGCACGTAGTCAATACAATTTAGCCTATTTTCTCCCATGTTACATTTGTT
AGTTGTACTGGTTTTTGGAAAAGGAAAAGTCATGCTGTTACAGTTCCACTAATAGAAAACAGATAATTTGGGAGGAAATT
AGATTGGAATAAAAGCACGTTGTTAAACAAGAAATCACAATAAAGTATTGAGTAGAGAGAAACCGTTAATGGAGACAGC
TATGTCATACCTAATATGCCCTCTTTTCATTGATCCTTAGAGGAAGCATAAGATCGCAGCTAAGTATGGTCTCTTGAGC
CAGCCTGCCATGATTCAAATTGAAACTCCATCATTCACCACTGTGTGACCTTGAGAAAGTTATGTAATCATCTTGAT
AGTGATTCTTACTGCTATGGACAGATTTGTGTTCCCCCAAATTCAAATGTTGAGGCAACTGTGTGAGGAGAGAGGGCC
TTTAGGAGGTAAAGTTAAATGAAGTCATAAATGTAGGGTCTTAATTCAATAGGATTAGTGGCCTTATAAGAAGGGGAAG
AGTTTTCTCTTTTTCTTCTCTACTTGCCTGCACCAGGGAAAGTCCTTGTGAGGACACAGTGAGAAGGTAGCTATCTGC
AACTGGAAAAGAGTCTTCACCAGAACCTGACCATGCTGACACACAGATCTCAGACTTGACGCTCTAGAAGTGTGAGA
AAATAAATGTTGTTTAACTACCCATTCTATGGTATTTTGTATGACAACCTAAGCTGACTAATAACTCACCTATAAAG
CAGGGATAATAACAGTGCCTATATGATAGTGTGTTTACAAGGATTTAATATATGTTAGTTTATATTATTTTATGTCCAT
ATATGTTAGCGTATATATTTATTTATGTCTATCTCTAATTCCTCATTGAGGATACTTAAGAATTCCTACTTTCCATGTTT
GACAGACCTGGTTTGGATTTTCAGTTCACCTCTTGTAACCTCTATTACCTTGAGCCAGTGACTAATGTATCTAAGCCTC
CATTTTCCCTATCTGTAAATGGGGATGATAACTAGTGCTGCTTGTCTCTTCAGGTTGTTGTGAGGATTAAAGGAGATATG
CATGACAATTCATCTGCCAGGTAGTAAGCATTCCAAATATGCTATTTACTGCCATCATTAGAGGTTTGCTGAGCTTCTCT
CTTTTGCATTAAAGTAAGAGACTATTCTCTCCAGAAAACCTTTGAACTACATGATGGAGGAACAAATAATAGCAGTTTCTC
CTCAACCTTGGAACTCTAAAAATGTTTTTTCTAAGCCATTCTTATCTTTATTTTGTTCATTAAAGATACATGCATTGTG
CATTTTGTCTTTTATTTAATATATGCCATGAGTGTCTTTCTCCCTATCATGATGTCAATGTTACATTACAGATTCTAAGG
AGCAGGGGCCATATCCCTTAAACATGATTTATTTAAATAACAATATAGGATTGGATGTGACTACTGCTTTTGCAATGA
AACTGAAAGATGGGAGAGTGAGATTTTCTCACAGCTATGGAGTGGCAGACCTGAGCACTAAAATCCAGTCTCAGAACCC
CAGTTATTATCTCACAATGTGAAGGCAGGAATCTATAGACAGATTATTGAACATCTCATGTATCATATCATGTATATCA

Fig. 9.144

Fig. 9.145

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GAAGAAGAAAAGTGTCTAGTATAGAGAAGTGAGACCTTTAAGAGGTGATTGGGTCATGAGGGTTCTGCTCTCAAGAATAG
ATTAATTGATTTCTGGATTAATAGGTTTTTCATGGGATTGGGACTGGAAGGAGAGGAAGAGAGACCTGAGCTGGCATGCT
CAGTCCCCCTTGCCATGTGAAACCTTGTGCCAACTTTGGACTCTGTGGAGGGTCCCCACCAGCAAGAAGGTGCTCACCAG
ATGCAGCCCCCTTGACCTTAGATTTCTTAGCCTCCGTAACCTGTAAGAAAGTTTCAGGTATTCTGTTATAAGCAACAGAAA
ATGGACTAGGCAGCATGTAAACCAAATTAATCAACATACACAAATTTTTGTAGCACTATTTTACTCTTTTATATAACAA
ATAAAGCCTTAAAACAAATACTATTTTGAAAGAACTTACATGAAATGGTCATAAGCATAATTTAAACTTTAAAATTAC
ATATGTAAAGTTTTTCATTGTCCAAACATGTTGATAGACAAAATGGAAAAGATAAATAGTTTGTACAGTTCTCTCTTCTT
GAAAAGTATTGCTTTGATTAAAATTCATATTGTCCCAACATTGTTTAATTCATAAATAGTAAAATAAATATGGAACATT
TTTTTCAATGCAGTATTTGTTTTCTTGAAGCTTTCAAAGGGAGAGACAAGAATCCCCAGAATATTAGAAATCTTTC
ATATTTTTTATTACAATCACTGTAATCCTGAATATTTTTTTCTAGCTTGAAGCCTAAAATTACCAGTACTTTAATATACA
AGATGAAAGTTTGTTTACTCATACCACTGCATATTATTGTTTGAACAATCATTTCTGGCAATAGTCATACCTCTGCGT
ATTATTGTTTCAAACAATCATTTCTTTCAATGACTCTAAATAAAAAAATTCCTGCTTCTCTTTAAAAAGACAGTGGGAAT
AAGTTTCTCTCCTTGAGCAATGGGGCCCCCTACTCTGCCTGGTTTTCTTCCCCCTTTCTTAGCACAGACACCACTGGAAGA
GCTGAGCCCTCATTTAAGATTGATTCTATATTAGTTTCTTAAGGATTCTTTAACAATGACTGTAAGCTGGGTGGTTTA
AAACAACAGAAATTTATTCTCTCACAGCTCTGGAGGTCAGAAGCCTAAAATCAAGGTGTTAGCAGGACTTGTGGTTCT
GGAGACTTTAGGAAAGAGTCCTTTCCTTGCCCTCTCTCCTTGCTTCTGTTGATTCCAGGCATTCTTTAACTTGTGGCATC
ATAACTCCAATCTGAGTTGATGGAAGTAGAGATCTCCATCAGTACATGGTCTTTTTCTCTCTCTGATTCTCTTCACTTC
TTCTGTTCTCTTCTAAGGGCACTTATCATTGGATTAAACTCCACCCTAATCCAGGATGATTTACCTTGAGATCCTTA
ATTGAATTACATCTGCAAAGATCCTTTTTTCAAATAAGTTTACATTTACAGGTTCCAGATGGACATATATGTTATGGGG
CCGGTTTTTTTTTTTTTTTTTTTTTTTTTGGAGACGGAGTCTCGCTCTGTTGCCAGACTGGAAGTGCAGTGGCACGATCT
CGGCTCACTGCAAGCTCCGCCCTCCGGGTTCACGCCATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCGA
TGGGGCCGATTTTAAATCCACTAACAGGTACCTTGAGACTAGGGCTCACTATACTTTGGTCAGAGATGCTCTAGGCATG
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GACAGCTAGAGCTTCTTGTTCATTAACCTTTGAAAGGAAGCATTTTCCCTGGGAAGAGGTGCCCAAGCCTCTTTTGTAA
GGTGGGCTTGTCTTAGGCAGTAACCTAAGCCTAGAGTCAGGTCTACTAATAATCCATTAAGATGCTGTGTTAGAAAAATC
CTCTCCAAATATGATGGAATAAGTATGTATAAGAAGTATTTTACAGATGAACACATTCCCACTTTATTAACCAAGTGC
CTGTGTGCTTGCTTAAATTTTTTACATTTTTCAGACACTTAACATATTATAGAAAATAACATCCAAAGAGAACATGATGT
TTAGCTTTTTTGGATTCTTTGGGATAAAGAGCAAACCAATTTTGAATTAAAAAAAAAAAGACTGTAGATACCACATCCTT
AAATAACTCAAAGTTTTTGTACCAAAGGCTACTTTTTTAAAGTTGAATCAAATGAAATAGGTGTGAACACCTATGTTTT
GAGAGGCAGAGAAATTGGTTTTTTTTTAAAAAGTGGGAATAAGTGAGTATTTTTTAAATTTGAAAAGGGTGTAGATTGGG
TTCTTCAGAGATTGCTAAAGCTGATATCATAAATTACCTAAGGACAGGGAGAATGAAGTGAATTGTGTAAATGTCTACA
ATATTATACTTTTCTTGCTTATACTAAGCTAACTAGTATGGATGACATGTAGTAAAAGTTATAAAAGGAAATTCATCT
TCACCTGTGTACATGCAAACTGCTGCATTTGCAGATTAAGTCAAATTTATATATAGAGAGAGACACACACACATAA
CACACACACACATATATTGAAATCTTTAAGTACATCTAATTTTTTTATGACTCAGAAAAGAATCCTTATTTAAGCCCTT
TTTACTACAGCAAAAAATGTTAATGTCCATTAATTAATGGACTTTGCTCTGATTTGGGGATGATAATTGCAGAAATGA
GTAGGGATATATATATGATCTAATATATAGAAATGGATATATAGAATATATATGTATATAGAATATATAGAAATTTCTA
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TATAAAGAGAGATATATACAGAGAGAGTAATAACAGGTAGTCCAGGAGTAGGACAGGAGATTAGTGGCCCAGAAATCAA
TACCATATTTGTTCTATTTTCATCTTTTTTGCAAGACAAAATAGATACCCAGGGGGCTGGGGAAGGTGACCAGTAAGTTAC
TTCCATTTATTTTCTTTTTCTTTTTCTTTTTTTTTGAGATGGAGTCTCACTGTGTTGCCAGGCTGGAGTGCAGAGGCACA
ATCTCGGCTCACTGCGACCTCCACCGCCTGGGTTCAGTGATTCTCCTGTCTCAGCCCCCGACTAGCTGGGATTATAG
GCGCGCACCACCACACCTGGCTAATTTTTGTATTTTCGGTAGAGACGGTGTCTCAGGATGTTGGCCAGGCTGGTCTCAA
ACTCCTGACCTCAGGTCATCTGCCACCTCAGCCTCCCCCTATTACTTTCTTCTTTATCCTTTTTCTACTGTTTAAAAGC
TCAAACTTGTGAACTCCCTCTCCATCAGATTCTGTGTGCCCAAAGAACTGTCTTGGAGGAGAGTTTCAAATGTCT
TTCATTTTAATGTCTGGCCTTGGTGATTGTTTAAGAAGTCCCCTGAATGAATTTTTTGGGTGAACTCGGGGCCTAGACTG
GCTCAGGAAATCTGGCACTGAGCCCCATATTTACTTTGGCCAAACACAATACTTGCTGTGCAGCCACATTCATATGCA
CTGCTCACACCACGCAGCTTGAGAGCTTTGTCCCTGATTCAAATCTGCTGGGTATTATTAGTCTGAAAATTTACTTTT
ACACCAAGCATATAAAACAAAATGAAATACAGTTTAAGAAATCAGCTCATAACATTTACAATTAAATTCATTAATCAAGG
CAGCTTATGGAATGCCACATGTGAACTGTAACTTTATAAATATTCAAGTAGTGAACAACTAGACAATCACATTGGCA
GCTTTGTTTCAGTATAAGTGATGTAGAATATTACCATCCATCAGCTTGCAGTTTTGTTTTAAACTTGTGTAACCATGT
ATATCATCAAAAAGAACTTTTGGTTACATTTCTGTTTTCCATATGTTTCCCATTTACCTTCTCTAGATTTATGTGCC
TTCGTACTAATTAGCACCTGCATGAATTCAGAAAGCAGCACTTGAATCAAACCAAGTCAAGCTGTCTGTGAAGGAAAA
AAATACCAAGTTCCAAGATTTCTTGTTAATGGAGATGAACAGTACACTTTGTTGATGTTCTTATTCACATGTGTCTTCT
TAAAGTATGAGAATTTAGTATATGTTCTTTTCAGTATATGCAAATATATCCACATGGGTATTTTCAAACCTATGCTGCAGT
GTGCTTTGTCTGGTTACCAGTTTTTATTCTAGTACAAGAATGCAGGATGTTATTCGTACATTCACACTATATTAATTT
AACCTCCTAGGTCTGGGAGGAGAGAGGAGGAAAATTAAGGAGGCCCAAGCTTTGCATTTGAGAGAAAAAGTGAAGTGG
GTCTTGAAAGATGTCTTAAGAAACAAAAGAGAGAAAATAGAGGAAATGTTAGAAGTTGAGGTAAAAGTGTGTCTCAT
GGAGTAAAGAAGAGCATGGAGAGAATACTATTAGTCAAGAGAACTGCAGCTCAAACACTCAAATGCAGAAGCATCATA
ACAAGCTAAATTCACCTCTTCTTTAGTTTCATGTGTTTTGGTCTATGTAAGCTGAAGACAACTCTCCTTTTTCCATG
CTCCCATACCTCATAGAATGTGTTTGAAGTATTCTCTCCTCTTCTATTTTTTCAGAATAAATTTGGGAAAATTTGGTATTA
CTTCTTTAAATGTTTGGTAAAATGCAGCAGTGAGGCCATCAGATCCTGGGCTTTTCTTTGCTGGGAGACTGTTTATTAT

Fig. 9.146

TACTTTGATCTCATTACTTACTATTGGTCTATTTGGATTTTGGACTTCATGGTTCAATCTTGGTAGGTTGTATGTGTCT
AGGAATTTATTTCATTTCTTCTAGCTTTTCCAATTTACTGCCATGTAGTTGCTTATAGTAGCCTCTAATGATACTTCTAA
TTTCAGCAGTATCAGTTGTAATGTACCCTTCTTCATCTCTGATTTTATTTATTTGTGTCTTCTTATTAAGATGCTATGT
TAGAAAAATCCTGTCAGAGTCTGGCTAAAGGTTTGTCAATTTTGTCTTCTTTTCAAAAAATCAACTTTTTATTTCACT
GATCCTTTGTATTGTTTCTTTCATTTCAATTTCAATTTACTTCTGCACTGATCTTGATCATTCTTTCTTCTGCTCGTT
TTTGGTTTGGTTTGTATCTTGTCTTTTCTAGTTCTTTCAGGTGCATCATTAGGGTTTGTGTTTGTGAAGTTTCTTCTT
TTTTTGATATAGGGTATAAACTGCCCTCTTATTACTGCTTGCAACTGTACCCGTTAGGTTTGGTATGTTTCTGTTTCC
ATAACCATTTGTTTCAAGAAATGTTTCAATTTCTTAAATTTCTTCATTGACTCACTGCTCATTGAGGAGCATATTGTTT
AATTTCTATGTGTTTGTGTTAGTTTCCAAAATTCCTCTTATTGATTTCTAGTTTGTCTATTGTGGTCAGAGAAGATAC
TTTCTATTATTTCAATTTTTTTTAAATGTTTAAAGACTTATTTTGTGGCCTAACATGTGGTCTATCCTTAAGACTGATTC
ATGTGCTGAGGAAAAAATATGTATTCTGCAGCCATTGATGAAATCTTCTGTAAATATCTATTAGTTCCATTTGGTCTA
TCATGCTGATCAAGTCCGATGTTTCTTGTGTAACCTCTGTCTAAATGATGTGTCCAATGCTGAAAGTGGAGTGTGAA
CTCTCCAGCTATTATTGTATTGGGGTCTACCTCTCTCTAGCTCTAATAATTTGCTTTATATATGTGGATGCTCTA
GTGTTGAGTGCATATATATTTACAATTCCTTATATCTTCTTGTGTAATTGACCTTTTATCCTTATATAGTGACCATGTA
GTCTCTTTTTTATAGTTTTTGTCTGAAATTTATTTTGTCTGATATAACTGTAACCTATTCCTGTTCTTTTTTGGTTTCCA
TTTACGTGGAATATATTTATCTATCCCTTTATTTTCAGTACATGTGTGTCTTTATCAGCGAAGTGCATTTCTTGTAGGC
AACAGATTGTTGGGTCCTGTTGTTTTATCCATTCAACTACTCTGTGTATTTACTGGGGAGTTTAGTCCATTGATATTCA
ATATTAGCATTGGTAAGAAAGAACTTACTCTGCCATTTTGTGTTTTCTGGTTGCTTTGTTTTCTTCTTTCTTTCTTCC
TTCCTTCTGTCTTCATTTTAGTAAAGGTGATTTTCTCTGCTAGTATGTTTTTAAATTTCTTGAATTTTATTTTTTGTGT
ATCAGTTGTATGTTTTTGTATTTGAGGTTACTACAAGGTTTAGAAATATCATATCATAACTGATTATTTTAAAGCTGACG
ACAAATTAAGAACTGATTGCATAAACAACAGACAAGGAAAGAGAAAAGTAATAATTTTACACTTTWACTTTGTCTCTCT
ACTTTCTACTTTTTTAACTTTCATTGTTTCTTTTTTTTTTGTAGATGCAGTCTCGCTCTGTTGCCCAGGCTGGAGTGCAGT
GGCGTGATCTTGGCTCACTGCAAGCTCTGCCTCCTGGGTTACGCCATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGA
CTACAGGCGCCACCACACCCAGCTAATTTTTTGTATTTTTTAGTAGAGACGGGGTTTACCATTGTTAGCCAGGATG
GTCTCAATCTCCTGACCTCGTGATCTGCCCCGCTTCAGCCTCCCAAAGTGCTGGGATTAGAGACATGAGCCACTATGCCT
GGCTGTAGTTATTATTTTTGATTGGTTTGTCTTTTAGTCTTTCTACTCAAGATATTAGTAGTTTAAACATCGCAATTAT
AATGTTATACTATCCTGCGTTTCTCTGCATATTTACTATCATCAGTGAGTTTTGTATTTTTTAAATTGATTCTTATTGCTC
ATTAATGTCCTTTTCTTAAAGAAATCCCTTTAGCATTCTTATAGGACAGGTCTGTTGTTGATGACATATCAACTTTTGC
TTGTCTCAGAAAGTCTTTATATCTCCATGCATGAAGGATATTTTTGCTGGATATACTATTCTAGAATAAGAGGTGAGAT
CTTTTTTTTTTTTTTTTCCACACTTTGTGTATGTACATACCCTCTCTGCTGTCAGACATTTTGGAGCGCCACTATATAT
TATTTGTTTATTTTCTCTGTTGCTTTTAGAGTCTTTCTTTATCTTTGACCTTTGGGAGTTTGATTATTAAATGCCTT
GAGGTAGTCTTCTTTGGGTTAAATCTGCTTGGTGTCTATAACCTTCTTGTACTTGAATATTGATATAATCCCTAGCT
TTGGGAAGTTCTCTGTTATTATCCCTTTGAATACACTTTTTATCCCTATGTCTCTCTCCACCTTCTCTTTAGGGCCAAT
AACTCTTAGATTTGCCCTTCAGAGGCTATTTTCTAGATCTTGAGCTTGTGTTTCATTTTTGTTTGTGTTTCTTTCT
TTTTGTCTCTTCTGACTGTGTACTTTCAAATAACCTGTCTTTAAGCTCACTAATTCTGTTTTCTCCTTGATAATTCTGC
CATTAAAGAGACTCTGATGCATGTAAATTACACTTTTCAACTACAGAATTTCTGCCTGATTCTTTTTTATTTCAATCTATT
TGTTAAATTTATCTGATAGCATTCTGACTTCTTTCTCTGTATTATCTTGAGTTTCACTGAGTTTTCTCAAACAGCTAT
TTTGAATTTTCTGTTTGTAGAGGACACATCTCTGTCTTTCCAGGATTGGTCACTGGAGCCTTATTTAATTTGTTTGGTGA
GATCACGTTTTCTTGGTGGTCTTGATGCTTGTGCGTGTTCATCAGTGTCTGGGCGTTGAAGAGTTAAGTATTTAGTGT
AGTCTTTGCAGCCTGGGCTTGTGTTGTACCTGCCTTTCTTGGGATGGTTTTTCCAGGTTTTCAAAGAGACTTGATATTGT
GATCTAAGTTTTTGTTCACTGCAGCTATGTCTGCATTAGGGGGCACCCTAAGACCAGTAATGCCATGGCTCTTGCTGAC
CCATGGAGGTGTCCCCGCTTGGTGTCTTGAATAAGATCCAGAAGAATCTCTGGATTACTAGGCAGAGATTCTTGTCC
TCTTCTTTTACTTTCTACCAACAAATGGGGTCTGTCTCTCTCTCTGTCTTGAGCTACCTGGAGCTTGGGGAGGGGT
GACACTAGCACCTTGTGGCCACCACCACTGGGATGGCGCTGGGTGAGACCTGAAGCCAACACAGCATTTGGATCTCTCT
CAAGGCTGCAATGACCAGTACCTGGCTACTGCATATGTTTGTCTCAAGGTCCTAAGGCTCTACAATCAGCAGGTAGCAT
AGCCAACAAGGCTTGTATCTGTCCCTTCAGGACATCAAGTTCCCTTTTGGCCCTAGGAGAATCCAGAGATGCCATATGGG
AGCCAGGACCTGAAGTGAGAAATCTTAAAGAGCTACCTGGTGTCTATTTTACGTGACTGACCTGGCACCAAGCTATA
AGACAAAGACCTCCCCGTTCTTCCCTCCCTTTTTCTAAAAGCAGAGGAGACTTTCCCTGAGGCTACCACCACCTAGACC
TATGGCAAGTGCTGCCTGGCTACTGCCAGTGTTTACTCAAGATCCAGTAGCTCTTCAGTCAGCTTGTGGTAAGTGAAAC
CAGGCTGAGATTCTCCCTTCAGAGCAGTGGGCTACCTTTTGTGCTGAGGGCAGGTTTCACTGCTGTCCGAGAGCCAG
GGCTGGAATGGGAGATCCCAAGGGCTGCTTGGTACTCTACCCTACTGTGGCTGAGCTGGAGCCTAAGCTAAAAGACA
AAGTCCTCTTTACCCTTCTTTCTCTTTTCTCAAGCAGAAGCAGTCTTTTCCCATAACTACTACAGCTGGGAATGTTCT
GGGTCACACCTGAAGCCAGCATTTCTCAGTCTCACCCAGGCCACAGTGAGTACCACCTGGCTATTGCTGCTGATTAT
TCAGGGGCCAAGGGCTCTTTAGTCAGCAGATGATGAATGCTGCCAGGACTGCTTCCCTTCCCTTCAAGGCAGCAGGTTCC
CTTTTGGTCCAGGGTATTTCTAAAAATGTCATCCAGGAGGTAAGGCTGTAAATGGTGGGCTCATGACTCTGCCTTTTGC
CCTATCCTACCATGGATAAGCTGGTATCCAAATTGCAAGACAAAGGCTCTACTCTTCCCTCTCCTCTCCTTAGGTGGA
GGGAAGGAGTCTCACCTGCAGCTGCAAGCTGCTCTGCCTAGGGTTGGGGGTGGGGTGGCACAAGCACTCCTTTGGCTGC
CCCTGCTAGTGTCTCACTAGGTCACATGTCCCCATAAACAATAATTTCTAAACCCATCCAGCATCAGTACTTGCCAG
GAATTTCAAGTCTTTGTGGCTTAGACTGTCTCAAGTTTATTTAGTACCCCAAGAACACTTTAACTTATGGCAGTGAGGC
TTGCTGGATTTCAAGATTCTGACTGCTGGAATGGGCAATTTCACTCTGGCTAGGGCTGGTCTATATTCTCCCTCCACTGG
CGTAGCTGAGTTCTGCCAGTGTGCTTTCCACTGTGACAGGGCAGCACTGAGTTCCAAAGCAAAGTCCCCCAGTCGC

Fig. 9.147

Fig. 9.148

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TTCATTAGTATCATGCTCTACATAAATTTGCTAGCCAGTCGTAGATGTCAAAGTAGAATTCAATCTATACATTAGTTTT
TCTCTTCCTCCTTTGACCTCCAGCCGTGTAATGAAAGACTTCTTCAAGAAGGATGGCCGGGCATGGTGGCTCACACCTG
TAATCCCAGCACTTTGGGAGCCCGAGGCAGATGGATCACTTGAGACCAATAGTTCAAGACCAGCCTGGCCAAAATGATG
AAACCCCTCTCTWCTAAAAAATAAAAAAAAAAAAAAAAAAAAAAAAAAATTAGCCGGGCATGGTGGTGCACCCCTGTGGTCC
CAGCTACTTGGGAGGCTGAGGCAGGAAAATTGCTTGAACCCAGGAGGTGGAGGTTGCAGTGAGGTGAGATTGCACCATT
GCACTCCAGCCTGGATGGTAGAGTGAGACTCTGTCTCAAAAAGAAAAAAGGAATCTATGTAGTTTTATAGATTTG
CAAAAATACTATATTCCCAATGTATGCTGTGACTCTTAAAAGTATTTACTAAATGTTGCCAATATCTAGAACCATCTC
TTAAAAAATATTTCAAACATATCAGTGTTAAACAATTAATGGTGAAACAACAATGTGATTAAATGGTGTTTTAAGTAGGT
TTTAAACCAAATTATATGTACATTGCATTGGCTTGTGCACTCTAGTTATTAAATGCATCGATTTTACCCTGGGAACA
CTCTTATTTCCCTACAGTTTACCTTTTAATAAGTGCAGTTGGCACATGTTTTGACCAGAGTTATTATATGTCTTATTGT
TCACTGTATGTCTGGGACTTTTTTTTTGTGTTTGCATGAACCTTGGCAGCAAATTCAAAACCTTTCTCACATTAATAAAGCT
ACTGGTTAAATATGTCAAGGGCAGAGTAGCTCATTCTTGTAATACGGTTGTATATACACACACCTTTGTTTTGTGTT
ATAGAAAATGCTAATTAGTATTTCCCTGATGACATGTGAGCTTGTTCATAGATTATTTTCCACTTACCTTAGGTCTTGG
TGTATTAGGCAGGCTGCGAAGCACAGCTTGCAGGGGAGGGCCACCATTGTCCTTTTCTCCTTCAAATATTATATAAAG
TTCAGGAAAAAAGCATTCTCATAGATTATATTTTGTGACTGACTACCCGTATTTTGGGGTTTGTGTGGTGTGATGAGT
AGAGATTAAATAATGGAACCTAAGTTTATAGATAGTTCTGGGGACTAGGAGATTAAAGACAATAACAGATGTGGGGA
AAAATTACTTTTGTAGAAATATTCAAGAGAAATTATTTTCAAGAGAAATAAATGTATGTTTCATGTTTAGTTCAATGTTAT
TTTCAAAGGATTATGTACAAGAGAACCCCTGTAATTTTTTCCGGAGTGATGATGTTCCCTGATAATTTATGTCACCAA
AAAGAAATGTGTTCTAAATAAATGAAAATTACCTTTAGAAAAATACCTGATTTACTCATACTTCTTATTAAGAGTAAGA
CACGAATATGAGAGGGGAAAAGTAATACAATCCTGACATAATGCAATGCCTTAGTCCCTAGTGAGGTAAAATATGTGATA
CATATATGTTTAAATGGACATTATTGTGAAGAATACAAGAACAATGACAGTTATTTATATTGCACTTAGAAAGGATACA
AATGAATGGTTTTCTTTCTCAAAACAACTGTAAACTAAGGCAAACCATTTTCCCTTCCAAGGTCTGCTGAAGCTGAC
AGCTGCCACAAAGCTTATTTCCAGTGTTCTCCAATATGTAGCCACAACAAGTTGTTGAGTCTCTTAAATAACTAATGT
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CAATTTCTTTTCTTTCTTTCTTTTCTTTTACACACATAC
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CCAGGCTCCTTACCTTTCCAGGCTCCTTAACCACCCCCCATCTTTGTTTTTCTCCACAGAACTTATTATCACCTGATA
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CACCTAGGACATGTAGTAGTTATTATATGTATTAAATAAATAAGTTATATGAAATATATCATTTAACAGCTGATTTGC
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TTGCCAGACTACTCTTTGATAAACCTGTCAATACAATTAATTTGTTGATAATATGCATATCTCATCACCAGTGCAGTC

Fig. 9.149

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TTACTTTCCTGCATCTGAAAGAAGAAAGACAAGAATGCTTTCTGCTTCCTCTTGACATTTTCCTTCCCGGGAAATATTT
TCCTCAAGTATTTAAGGTGAGTTACATTAAAATATWGAAAAATTAGCTTAAATCTTTTCTTATCAGAAAAATAAGCTGA
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GATTCTGGTTCAGCAAGTCTGGAGAGGACGCATAATCTCCCTGTTAAGGGTTAGATTCTAAATAAGTGGTCTTCAGGCC
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TGTAAGAGAATAGCTCTTGCTTTGGAGTAACTTCTCAGTCAGAACTAGGTCTTGAATTCAGACCAGCTCTCATCT
CCTTGGTCCACTCTAGGGCTGATGGAGGCTGCTGAAATCATCTTTTGCTACATTAATACTCATTTTTTCGCTCAAAGATG
ACAAGGAGTATTTAAACGCAGAGAGAATGTTATTTATAAATGATGATTGTTTCGTGATAGAACTTGCCACATAGTATTT

Fig. 9.150

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TCTCTTTTTTAACTTTTATTTTAGGTTTCAGGGGTACATGTGCAGGTTTGTATATAGGTAAGCTAGTGTGATGGAGGTT
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TGTAATTTCTGCAAAGTCTTATCATGTATATTTTGTCACTAGTTTATTTTACAGATGAGGAACTAAGGCTCAGCAAAG
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TAAAGTCAGATATTTGAGAAATAAAATTCATTTCTCACTCCAAGAAAAGTTTCTGCATATCCAAGGATGTGGGGGATA
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TGTGCTACAACAGTTTTTAAAGTCTGCCAGTTATTTAATCTGGGGAGATTACAAATACAGTGTTGACGACTGGCCTGC

Fig. 9.151

Fig. 9.152

Fig. 9.153

Fig. 9.154

CATTTTGATTCTAAAATGATGTACCTTCTGTGTTTCATGGAAATGGGAGGTGGTGGGCAGAGGAGGTAGAAGGGAAAGG
ATAGGAACAAATCCCCAAGCAGGGAATGTTCTAAGTCCTTGTCTTATTCTAATTGGATCATCTGGGTTTCAGTTGCCCA
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GTAAATATATTGCTATTAACCTCTGAGAGTTAAAGAAATATTTCTCTGTGTGTTTATAAAAAAGATGCTATGTGAT
ATGGTGTGTGAGTTTCAACATTTTTTCTATACCTTCAATAAGTTTGAAATAGATATAAATGTTTCTTATAACTGTGACT

Fig. 9.155

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GATTCTTATAAACTACACATTGATGGCCTGTTGTCTAAGTTCCATTTCATTGAAAATCTATCAGCACTGGAGCCCCAGT
TAGCATTCTTAACAGGGCATTCTCTTTAAGACTGAACACTAAGTGTGCACCATTAAAAGAAGCTGCATTCTTCAACTTG
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Fig. 9.156

GATAGATCCTGTCGACTGACTGACTTAATGCTATTAGATAAAATAATTTTTTAAATTACATGTATTTCCAAAAAGTACAA
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ATTTTATTGACACAATTTTACTTGACTTTGCAACCTGACCGTTTTAGATACCTTTTCTTCCAGTCTTTTCTAAAAGAT
CAATTGATTAATTGGGTGACTCGATGATACAAGCACACGATCTTCAGTATCAAAGATACTCATGACCAGTAAACCCC
TCTTTAGTTCTCAATTATCCAATTCCCTTTCCCTGGAGGCAACTGCTACTCTTGTATTTCTTAATGGAGATCTTTTATGT
GTTTACTCAAATGCTAAGACATAATATATCATATTCTGCTTCTGCTTATCTTCCCTAATGATATAGCAAGGAGATTGA
TTCCTAACTCTGCATGCAAATCTGCCATAGTGCTTTTAGCAATTGCAGTATCGTCCATGATCTGGATGAGACATAATTT
CACCAGACTCCTGTTGGAGTCTGTTGGATTGTTTCCCATCCTTTGTCACTACCTACAGTGTAGTGATGACTATTCTTTT
ACTTATTTGTTTCATGTAAAGTAAATATATCTCTAGAATTAATATTTGGAGGTGGAATCCCAAGTCAAAGGAAGGAAAT
ACATATTTTTTAAAAGGATTTTAGTAGATTGTTCTGAAAAAATTACACCAAACCTTACCCTCACATGACTATTTTTTGCT
TTTTCTCAGTATACTTTTTAATAGATTGCATTATAAAATTTTAATATTTACCATTAAATAAGTGAAAGCAATGTCTAA
TTGCAGTTTTTAATTTTATTTTATTTAAGGAAGTTAATCTTTTTTTTTTAGCTGAAGTCGAAAAATATGTTTCATCTCTTT
TTCATTGAAGCTCATTAGTTTCCAAAAATGTGTCCTCATTTTCATACAAAGTATCTTTGAACCATTGCCAGGTCATAGAA
GTGACTCTTGAAGATCTTCTTCTCTGGGATCAAGGGTCACATCTCGTTCTCCTGGCTTCATACCCCTTCTCTCTCTACAC
ATCTCAATTAAGTAAGCTTCCCTTCTTATCCTCTGAAATTATAATCATCTTTAGGCTCCAGATATAGGTGACCTGTGC
CATCTAGTCTGACTTCCAGTTTGCAAATATCTTCAAAAAGTGGAACAGCTTTTACTTTCTACTTTATAAATTCATAC
GTGCTCAGGAGGACTTTTATTAAGATTGGGACATATTGCTTAGTTAGTCTCTAGGGTGGAATCATAATTAATGAAAGAA
CTCTGTCTGATGGAAGGCTTAGTCTAACAAATCATCCTAAACCTTCATTAGACATTGTGTAATCCCATGGACACTGTT
GAAGAGACACTTGATTAGGCAGATCGTGACAACTTCAGCTTTGCCCATCATACATCCTCCAGAGACTCCTCAAACCTCAA
GACAGACTTTCTACAGGAAGGAATGGCTGTTTCTATTTTCAGAAATATTACTGATACCAGAGCTTTGCTGCATCTTCATGT
ACTTTCCTTTTATTTTTTATTTTTTATTTTTTTTACAGACAAAATCTTGCTCTGTTGCCAGGCTGGAGTGCAGTGGCATCAT
CATAGTTCAGTGTACCTCAAACCTCCTGGGCTCAAGTGATCCTCCTGCCCGGCTTCCGGAGTGTCTAGGATTACAGGA
GTGAGCCACCACACCCAGTCTTCATGTACTTTCTTATGTAAATTTATTTTGTGTTGAGATTTTCTTTGTGATGGAATAAG
CAATTCCTTCCCTTAACCTCATTTCCCGAGACATTTTTGGAATTTGCAATATATATGAAAAAGCACTTATTTCTCAGTGATG
GAAACACCATTATTTCTTTTACTAGAATGTGATCTTCTGGTTTTAATATTTATGTCTCTGTGCCAAGGTGAAAATATTG
TAGAAATGAAAGTATAATGAGAGAGAATCCAGATAATAAGATTTTCAATAAACAAAAAAGAGTATGTATAGGAGTCC
TGTCTGCTGCAAATTTTAAGAATGGTTTTCTGTCAACATACAGTTGTCTTACTGAAAACCTTCTATAAAAAACAA
GACACAAATGATTGAGAAGATAGTAGAGACTGAAACTAGAACTGTTGATTTAATAAGCTTTCTTACTCATATACTTAAT
CCACAGTAAATTATTAGCAAATTACATTTTCATAGTATTTTCAGAAATTAAGATCATAGTCAGGGCTTAAAGGAATCCA
GTTGTTTTTAACAGTGTGAGGCAAGCGTTCTAGCAGCAATCACCTGTCAGAAATGACTTAGGTTGGATTGTTTGCACC
ACACACTTTACTATAGTTACATGTCTTTCATGTCCCTTTGTATCTCTGACTCAGTTTCTCATCAGTGGTAATAGCAGA
ACTTACATTTTAGTGGGAGGATTAAAATAATGTACATGAAATCAAATATATAAGTCAGCACTCAATAAATAATAGTTTG
TCATCATTTATTACTATAGTAAATGACCCTATCTGAAATGGGATATAAAAAGGAAAGTGAAATTACTAAATAAATATGA
GCGTACAAAGACCTGACAATTTAGATACAATATACTCACTCTGTCTCAAATAGTGTCTTCTGAAGCTTTAGTCAGAAAT
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TCTACAAAAGGCTGTGGCAGTGGGTCCCTGAGTAAACCACCAGAAGAAACCTAAGGGGCACCTCTGTTTTTTCAGTTAAT
CAGTTTTTAGCTCAACTGGGATTTAATCCTGAAAATCTAAATTACATTTACAGACTTTAAGTTATTGTGAATATTTACAC
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AGAAAGAGTCTTAGGTGACCTGGTGTGGGAATTGCTAAATCTCAGGCTACTCACTGAGACTGTCTGCTGCTGTGCTGGG
CACTGCAGTCCCTCCAGAGCAGGCTGTGAGCATCCTACCGTGCTCTGAGTAGGTCCACTGGGGAACCTGGTGGAGTGAC
ATCATGTATCAGCCTCTAGCAGCAACTCCTCTCTGTAGCCCCATGTCTTCTGTGGTACAAAGGAACAAGTGCTACAG
GGCATGTCTGTAATCGTTATCCCAAGGGCACGCTCCACAGGCATCTAAGGGTGAAAGGTACTGCCAGTCTGTGTTGGGTTT
GTTATTTCCATAGAGCTACACAGGAAATAACACCACCAAAAATAACACATTCAAACCTCAGAGGGCAATCTTCCCTAACT
ATTCATAGGCACACGTCAGGCATTCTATACATATAACCAGCTCTTTGCTAAGCATAGTGAAATGCCCTTTTACATTGCA
ATTAATTATTAGCCAACATTGAATAATTATTGGTGAGAGGGTAAAGGGCAAGTGAAATAAAAAATAGAGCTGGTTTATTT
TTAGGAAGACACTATTTTAAATGTGTTGATTAATCAGACAGGTGTTTAAAGCATTGTGTTAGAGTCAATTCACAGAAAT
CTCTTTTACATGCAGGTTACAGCCAAAGAAAGAAATAATAGCCAACACATTTACGATTTCAATTGCAAAAATTTGTCATA
TTTTTGAACCTGTGTGGGTATTTGAAAATATTAGCTCCTACTAGGTCCAGTTAAAGCTTTTAAATTCTATAAGGTTTCAG

Fig. 9.159

Fig. 9.160

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CTTGTCAGGAGAAGGAAATGGTAGAAATCAGATGAATGGTGAAGTATGACCATTAGAAACCACCCTCTATACACCTACA
CTGCCTATTTTTTACATCAAACATAGCGAACTGTCTTTAGGAAGCAGAAAAGGCAGAGCTGAGGAAATGAGAGGCACCT
CTCATGTCTTTAAGAAAACAACCTAAGTGGCAATTTCTGGAAAATAGGTTATTGTTTGTGTCAGGTATTAGCCTGAAATTTT
GCTTGAATTGTTTCTCCCTAGGAAATGAAAGATGTTCACTTCTTAGTTGTTTGGAAAAACAATACTTACTAAAGATT
ATAATGGCTATCAATTTAAACAAACAGATGCTTTTAGTGTCTTTGTGAATAACACTAGTGAATTTGCAAGAATTATTGC
TTTACCAACAAAGCATAAATTCCCATGTTGCTTCAGTACTTGTGCAGTGTGTTGTTCTGGAGCCTCTATTCAATATCAAC
CAGCGTATAGTAGTTCCCTCCGTGAGAGTCTCTTGTA AAAA CTTACATTCACTTCTTGTATTATTTTATCTGAGTGAATA
TTTGACAAAGACCAAAGAAATCCCCCTTGAGACTGGTACCCATTTTCTCTAATTTGAAAGATATTGACTAATTTTGTTAG
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AATCTGAGGGGCAACTGGTGTGTTGTTGATCTATGTTGAACTTATGGATAGTTATGTGAGTGTGAGATAGAATCAGACA
TCTATGTGAGCTGGCGCATCTCCATTCCACATTCTCTCTCATCTTTCTTCTGGGACTGATGAGCTTGCCTGGGTGTGTT
CTTCTCGTGGAGATGGAAGAGGTGGAAGAAAACAAGCCCAACACAAAAGCACATTTTCAAGTCTCTGTGATGACAGTTA
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CATAGATGCAAGGAGTAGGGATTTGGGGCCAGATGATACATCCTACCACAGAGAGGTAGGATACAGGGAAATTCTAGGC
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CTTCAGCTGCTGCCCAAGTGTCTAAGAACACTCATTACCTCTCAGATACTCACATACCTATGAAATGCACCAATTTCA
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CCTTCATCTCTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCT
CCCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCTTCCT
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GCATCAGCATTGCCAGGGAGCTGGTTAGAAGGGAGAATCTCAGGCCCTTCTAGACTTACTGGATCAGAGCCCGCATTT
TGGCAACATTACCAGGAACCTCATATACACACTCAAATTTGCCAGGCTCTGAACTCTATAATCTCCAGCATTCTGATTTCT
TCCAAGTCCCTCCTTTCTCATCTTTGAAACAGAGTAATTTCTACCTCACAAGTCTAGTCGAGAATTAAATACAAACATG
TAAAGCTGCTGTGTTCCGTGCCTGGCATAACATGGCTGTCCCCCATTCCTGCCAACACACACCTTCAAGCAAGACAACCTT
GAGAGAGGCTGAGTGGGAGAGAGGGTGAATAAATGTTAATAACTCACATTAGAGTTGGCATTTTTTTTTCTTTTACTTG
TCTCTATTATATAGGGTGACTGTATTAATTGTGCAATACCTTTGCTTTTAAATAATACTACTGGGATAAAAGTGAGATT
CTAATCTCACATATTATTTCAAGATAAGATTCAGATACAGAGTTTTTAAAGAACCACCTATAAAAATCAAATCCACATAA

Fig. 9.161

ATTCAGAGGAAAAATATCCTCAATGCTTATATAATATTGGATGTTTAGGGTGATACCCTCTAAGCACAACCTTCAAAGGAA
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AATTA AAAAGGAAGTCACAAATTGAAAAAAATTATATGCATAGACAAAGAGTTGATAAATTTAACATAGAAACAGGTTT
TACAAGACAATAAAAGGATAAATGTAATGATTTTCAATAGATATAAGACATTAATAGGTGGGCATGTAAAAAATGTCAA
GATAACATGAAAAAATTAGTTTTACTACTCATAAAAGACATGCAAGTGAAAACAACCAAGATGCCATCTCCAGTCTATC
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TTGCAACAATATTTATGGGAAATTCTTTGGCTATATCTATCACTATGAATTTATGCATACATTTTTCACCCAATAATTA
TACTTTTGAGAAGTTATCCTAAGGAATTAATCTAAAACCTGAAATGATACATGTTCAAAATTTTCATTAAAGAAAACAC
CAAACCTGTACCATAATGGGAACTATTCATAAATATGTAAATTATGTCCATATGATAGAATACTAGGCACCCATTAAATC
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CTAGTTAAATTAGACGAAAATATAAAATTTTAATGGCAGAAATATTTCAATTTATAACATATGCATAGGAACTGAAAA
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AGAGTCGAAGTTACACTTCTGTATGGCATTTAATTCCTCCTCCCAGAATACAGCCACTGTTCTACAGTACAGAGGAGT
CCTTCATATCACGATTTTTCATTGTGTCTAAGAGGCATGTGTTTCCACTTTGTCAATTATTTGGTCTAAAAGGATTTTTCT
TCTATTTTGCTGCTGTTTTTGTACAAGACATATAAAGAAAGGTGTTAAGGGAGTGTTAAATATAATCAAGGTGTTAAGG
AGAAGAGCCCAGAGAGAGAACTCTTAATCTTTTCATCTGGTCCAGTTGTCAAAGTCTTTTGTCTTTAGCCATCACTTTT
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AAAATACTACTGAACGACTCTTPTTTTTTTTTATTCTGTCTCTTTTTCTGTTGCAAACTGATTGTCCACTATAACCTGG
AGAGGGGACTCAATTTTTAGAACTGTCTTATACACCTCTTAATTAGTTACCTATTTGCTGCATATACCCAAAATTTAGC
AGCTTAAAACAATAATGAAATTTTATTATGCCACAGTTTCTTTAGGTCTGTAGTAATTTAAGAGAACTTAGCTGGGT
GATTATAGCTGACTCTCTCATGATATTACAGTCATATACATCAGGACTGCATTCTTCTGAATACTTGACTAGAGCTGGA
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GGGTGGTCTTCTCCATGGGGCAGCTTAAGGATCTTCATGGTTTGTGGGCTGTTTCACAAAGTGTGCGTGATCCATAGGA
GATTAAGGTGGAAGCTGCAATGTCTTTTATGATAGAGCCTCAAAGTCAACAATGTCAATCCCAACGTATCCTTTTTT
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CATGCAAAACATACTCCCTTTATTGTGGTTCCCAAGAGTCTCATTCTATTATGGCATCAGCTCGAAGTCCAAGATCTTA
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GGAAATTACACAAGGCCTGAACATGAGGAGATGGGAATCTTAGAATCTGTCTAACATAACCAGGTCTATGTGCAAATAAG
ATCAATCTCTGTTAGTTTACTCTTATGGGGTAATTAATTACCCCAAACTTAGTGCTTAAACAATAGCATAACTAAC
TTCTCAGTTTTCTATGGGTCAAGAATTCAGAAGCATCTTTCTTCAGTGGTTATGGCTTTGGCACTCATATGAAGTTGCAG
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CCAACCTTCAAAGTTACACATCATTTGCTTCTGCCCTATTCTGTTGGGCAATCCTGATGCAATATAGGAGGCACCATGAA
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GAAAGAGATTTCTGGAATATTTCCATTCTCTGTATGCCCACTGCCATAGTCTACTCAAGTTCTCATCTCCTATTGGAC
CTCTGCGGTAGCTTTATAAGTCTCTGCTTGTTCCTTCCAATGTCTATCCCTTTCAAACGCTTCAAATAGCCTCCTTAAAGC
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AGCAATTTACGTCCTCTATGCCTTTTTCTTTTCTGTTCTCTTTGCCCTGAAATAAGAACGACAGATAATAGTGACTATGC
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GAAAAGTTTGGAGGTTCTGGCGTAACCTCTTTGTTATTCTCTATGCAACTAGCTTAAAAATAACAACCTTTGATTATTC
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GTCAGTATGTACGTGTTAAAGAGTAGAGGGAGGAAGTTGAAAATCTGACATTTGATTTATTTCTCCTGGGACAAAGGAA
GTTTGGATATGTCAAACCTGCTGGCTAATCAACACGTTAATCAATGCAATCTCTAGTATGGATATAGTTATATTTTGCC
CCCCCTCTTCTTTCCAACTTCTTTCCAGTCTCATGACCAGCTTATTGGCTTCATATTACCAAGTAAATATATCAACTG
CCCATATGTATAGTACCCAGACAATGAGAACACTGCCATAAATATTGTGTTTTCAATACTACTTGCTTTTAGCAACAAT
CATAGCACCCAGCCTAACTTTAAAAAATTCATGTCTATCCAACCTTCAAACCTATAACAAAAAATTGAGTTTTACAGAAGAGT

Fig. 9.162

TTCAAAACATTTTTATGCATTAGACACTTCAGAAATTATCTCTTAAAAACAAAACAAAATAAATAAGGGACAAAACAA
TAGAAAACGTCAGGCACCAAACCTGTGAACCTCTCTTTTGTCCATGTATGATTTAAAAAATAAACTCACTGATGGTAAT
TTAAGAAATTTGACCTTTAGGAAAATCAGGCTGAGGCTCATCTTTGTACTTTTTTAAACAAGTCAATAAACATATAAA
TAATAGAGATGAGTGCCGCTCACAGGGGACACTCATATTTTCTAGGAGCCTGTGGATATAACAATGAATTTTCATGAAC
ACAGAGCACATTTCTACATGATCTCCTTTTCTTTACAAGTTATTTATGTTTTACTGGAAATATTAAACTTTCATGTCTC
TTTGTGTGCTCTTGTTTTCTTTAAATTTTTTATTTTTATCATCAGTCATTGTTCAATAACTCAAATCATATAAAATAAAAC
AGTTTCAACATCATTTCCGTACTCAAGATGATCTCTATTAAATAATGATTTTTTTTCTTGAAAAAGCATTTCTATAATTTTT
TTTAGTTTTCCAGATGACTATTTAAACCAAATGTTTAAATGAATGATTAATGAGTCTTTTATCTTATTTATCCCACCTG
CCTACTGCAAAAAGACATGAATTTAATACATTTTAGATTTTAAACAGTTGTATATCTGTAAAAAACAATTTTTACAGG
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TATAATGCTCACCAATGGATTTAGAAATATCCCATTTGTAATCATGAAGATCTCAATTATATGTTATGAATAAGAAACA
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GGCGCAGGCCAATGGTTTTAGGCAGGACTTTTACAGAGCAGACTTCTGTCAAATTTGCCATATGGAGTTACTGGGCTCAGA
TTATAGAGGTTTTCTGTTGTTTGCCTAATTTGGGAATTATCAGTGCTCTCCAGGATTTTCTCCAAAAACAAAGAACA
TGTTTCAAGAAACTGACACTTCAGGAAGAAAAGTAGTTTGATATTTGAGAGGTTTAGATGGTTTTCTAATATTTCTAAA
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TCCATCTCTCACCCATGGAGTTCATATTTCTGTCTCATGTCCACCCTCTGTCAACGAAATAGGCTCTGCCATTTCTC
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AATTCAGTGGGGTCTCCATGTGACGTTTTCAGCCTTTGCACCCTACTTGTCCGCTGCAGGCCAGGAGCTCTCTACTCCT
CATATGCATTACAGGACCCCATGCACATTGAAGATGTTTTCTCAGGCAGAGCAGACACCAGGCAAATATCTTCTGGTCT
TAGGAGCAAGTCTTCTGAGAATCAATGATGACAAATATCTTAAATGGTCTACTCGATGGTTCTGAAAATGTAAGAGCTA
AACTACCTGTATCAAAATCATCTTGGGGGCTTATTTAAACATATATGATAAGGCCCCACCTCAAACCTAAGGAGTCAG
AATTTCTACGGAAAGGCTTGGGAATAGGAACATTAAACACAAGCATCCCGGGTGGTTTTTATGAAACAAAGCTTGAGAAT
TATTGAATTCTTCTCCGATCTGACAGCTCTTCTCCTGGGCAGACCCAATTAGAGGGAACTGATGAGTTTTAGGAAATAC
TGTAATTCTAACTCATCAGTCCAGCAGTGCTGAAAATGCAGAATTTTGTCTTATCAAAGCCAATTAATGCTAACCTT
TTCAGTGTGTGTCCAGTACCACCATCGGTAAGGTAAATAAACAGGCAGTTTTTGCATATAATCTAGTGATTTTTTAATTTA
GTTTTTTTCCCTTACTCATCTTAAAAAAGCAGGGAGTCATTGGGCTTTTGTATCTATTTTAGCTGAGTTATTATTTTAA
CCCCTATCTTTAGTGCAGAATTAAATATCACCCCTTCTATTTGCTTTTATTCCTAATATCTGAATGGTTTCAGCTTGAC
GCAATTAGCATGTAGAGCATGCTCACCTACTGTAATCTCCAGATGAGTCACTCAGCTCCAGCACATAACTGCTTCTGG
GATCCTGTTGGCTAACCAGGCTTTGAAACCCAGGGAAAAGAACTGTGTGTTTTCTTCAGAAAGTAATCCATAAATGAGAA
ATTACCATAAATAGGCAGTTTTTATTCATCTAAGTGATTTTTTACATAAAAACCTGACAAGGCCGTAAATTTGTATATAGAA
TCAGGTTCCATATCCACACAGCTAATGCACACATGAATTATTCGATGAAATTTGCTTCAGTAGTATTATCTGAAATTGG
TTTCAGACAATACATTAAATGTTACACATTATGGTAAATTTACTGTATTTTTTGTTCAGCCACGTCTTGTGTTGGCAAA
TCTGCTAAATTTAAGTTTGCATTGTATTCACATGTGTTGCTATTTTCTGAGATGTCCCAATATATTTTAGCCATCAAAC
AAATAGAACTTCACCTCTCTGAGGCTCAGTATCCACACTTGTTTATCATGCAAGAGACATGCAGGTGAGAAAATAGGCA
ACTGCCATTCTATCTGTGATTGCTGTTTCATGGAAGAAGTAAAAATGAGCTGGGACAATATATAGGTGGCGCCAGGGATG
GCTCCTTGAAGGAAGTGATCTCACAGCTCAGACCTGAAAGCTGAGTGGAAGTTACCAGAGGAAAAGAGAAAGGGAATGC
ATTGGTTTTCTATTGCTGGGTCAGAAACTACCACTAACTCAGTGGCTTAAAGCACCATTCAATTATCTCATGTTTTCCCT
GGGTGAGGAGTCAGGTGCGTATCAGAGTCTCAAGAGGCTGACATCAGGTGTTTGCAGATGGCATTCAATGCTGCTGGGCTC
AGGGTCTCTTTCAAGCTCCTTCAAGTTGTTGGCAGAATTTAGTTGCTGTGGTTTTAGGACCGAGGTTCCCACTTTCTT
GCTGATCACCAGCCTGGGGCAGCTCTCAGCCCTTGAGGCCACCTGCAGTTCCTCATCCTTTGGCAGATCCTCTCACAA

Fig. 9.163

CGTGGCCACTTACTTCAAGACCAGCAAGAGAATGTTCTCTTCAGGAAGGGCCCAACTCCTCCTTATGGATGTTCTTCT
GATTCAGTCAGGCCCACCCAAAATAATTGCCATTTTTTGGTTAACTCAAAAATCAACTGATTTGAGATCCTTAATTACAT
CAGCAAAATCCCTCCACCTTCTCCATACATAATAACCTAATCATGGGAGTGACTGTTTCATCCTGTTCTGATTAGGCTC
ACATATAAGAGAAGGAGATCATATAGGACATGCATGTAGAGGATAGAATCTTGGGGGCCATCTTAAAATTGTGCCTACC
ACAAGGAAGGATTTTCCAGGCAGGAAGAACCAGATATATGAAGGAGGCCTAAATGACAGGATAGCAATTTCCAGTGTCT
GAGGCTAAGGCTGGAATGCTGACCTGGCCAGATCATGCACTGCTAGGTTTAAATGGCCTATGCATACTCTATTATGCAAT
TATTAAAATACCTGTTTACAAAGAATATTTGAAATATTA AAAAATGGAAAATGCATACCGTAAAATATTAATGGGAA
AATCAGGATATGTGTTTATATAACATAGTATCAGTTTTTGGAAATATATATGCATATCTTAAAGACTAGAAGTAAATAC
ATCCATCTGTTAGCCATGGATATCTCTGGGTGAGAAAGTATGGATGAGTTAGAGATTCTTTGTCCCTTTTCTTAGTTCT
AAAATTTGCCACAATAAATATGTATTCTCTTATAATAATAATTAATTA AAAACCTATTTCATGGTACTTGCAAAAACAAA
TGAATTTGAGCAGTCTCTTTGAAGGAATCAAATTTTGTACTCCAGGTGTTTGAGAGGTTGAAAAACAAGCTCAGGAATT
AGCTGCGCATGATCTGTACAAACAAGGGGATCAGTCAGTGGAATGAAGTCCAGCTTCAGCCAAGGGTGGTGAGTATAA
CAGACCTACTCTGATATGTTTCATCACTAAACATTGAGCAAGAAATCTAGTGTTTAAATTATGGAGGCATTAAAGCAAT
TAACCTCCCTCTACCTCAATTTTCTTATATCTAAAAATTGAGGGGAGCAAAATTCATTCTTGTACAGGTCTCAATTG
CAAGTAGCATGCAAGCAAGTAGGCTGAACTCTTGTCTGTCAATCTAGCTCCTTCCTGGATGGGGTATTTAATCAGCTCT
CCTTCCCAAGCACCCAGTGGACTTGAAGCTCACAGTTAGACAGAGATGGCTAGAGGTGCCAAGTGCTGTTTATCATAGG
TGACATCACCTGTGTGCTCAGAGTTATAAACATAGGCAGCTTGCCCTACTGCCAAATCCATTACAACCTTTTCAGAGGTGTT
CACTTAGGAAGAGGAGGAAAGAGTTTTTTTGAGAACCAGATGGGAACCTTAGAAATAAGTAACATTGACAACCAAATCCTT
CAGACACATCTTCAAGTTTCAAGAGAGCAGTTAACAAAAAAGAAAAATGTATTTTTTAGAAATTGTTTGGGAGGTGAAAG
GATTTTTCTGGGGGGGAGTAGGGTAGGGAATGGATAGAGATGATTTTTTTTTTGGCTCTGCAATATCTCAGAGATCTTTG
AGTTGAAATATAACAAAATGTATTTCTCTCTCTTGCTCCAAAACTGAAAAGCAAGTGCACTAGAAAATGTCAGAGTT
GGCCACTGAACAGACCATATTTCTAATTCATGGCAAGATGGCTAAAAATAAAAATTCCTTGAAATACGTATTTTAAATTA
GATTTTGACCTTTGATAGGGCTTATCATGCTATTTATATTTGTAGCAGAAGGAACCTGATTTTTTTAGTAGTTGCCTCATCA
AAGACATGACTCACATTTCTTAAGCATTACACCATATGTTTCTTGAGGTAAAAAATAAATAAATAAATTTTATATT
TTTTATACAATTAGTGAAATGTGAGCAAAAAATATAAGAAATCATTTAAAAAATATTTTTGGGGAGGATGTGGTCTAGAA
ATGTTGTTTTCTAGAAATTGAAATTATGTTAGCTATTTGGCCAGTGTAGCCATTAGGTACATTTATGAATATAGATGAG
AAGTAATCATTGTGATGCTAGTTTTATCATTAAGGAAAAAAGCAGAATTCAAAAGTTTAGCAATGAAACACTGGTACTA
CTGAAACTCACTTTATATTTGAAAATGTGAATGTGAGTTTAAAAAAGCTTTTGTTAGACTAACTGAAAAGAATCATTTT
AACATTTAAAAAGAAAAATATTATTTTAAGTGACATTTGGTGACATTATTATATGTGTACAAGCTGTGACTATAAGATA
ATGAGACAAATTATTATGTGGCCAGTGGAAATTAAATTTATCTTTATGATTACTAGCAACAGGGACTATTTTATTCAAC
TATCTTTACATAATAACAAAATTATTGTAAAATTTCTAGGAAATTCATATTCACCTTAAATATAGAGGAAGTACATTTGG
AAAAAATATTAACATATGAAAATCAGCATCCCAGTTAAGGAATCCCCAGTTTCCCACTTGGTGCTCAAATCACTTCACG
TTTTATTTCAATTGGTACTTTTTCTAGGGCAGTAGATTTAGGAATAGAAAACAGACTCTACACATGGAATACACTAAAGT
CTTGAGGAGGTGTGATATTTCCCTAAGTTTGGATAGGTGGTATCATCCTTTAGTCACTCTTGACTCTGCATTAGACTC
TCTACCTAATGATCCCCAAGTCATFTCACTTCTGCCTTCTTAACATTTTGAAGATGTCCTCTAAAATTTCAAGTCCCAC
CATCTCTGCCTCCGAACCTCACCTCCCTGACTTTTCACTCTTGAAGCACCTTAACAACCCCTCCCCTCCTGTGCATGTTCC
ACCTGGAAACCAGGGTGATCACTCTGAAATGCCAATCTGATCAGTCATTTCCCTACTTTTGAACCTTCAATAACTTCCT
AGCACCTATTCTAGAAACCATTCCAACATATCACCTAGGCAAAGGAAACCCCTTTTCTTACCTGGCTCCTACTTATCCC
TCCAGATTACCTTTTGACACCTCCCACTTCATCTCTCTGCAGCCCCCTTCTGCACACACACGGAGTTTCAGTCTTGC
CAAACCTCTTCGCAGCTTCTTGAGTGGACCTTGCTCTCTGCACTACAATCTCTTTCTGCCTAGAGAGTCCTTTACCTCCC
CCAATCTGGTTAACTTCTACTGTTTTATTTAAGTCTTAACCTCTAAGTAGATGTCATTTTTTGAGGATGCTCTCTCTGTTCC
CACTTTGGAACCTTGGGCTTCTCTGTTGTCTGGGTGTCCCGTGATTCTCTGTGCATAGCCCTCATCTCATTGAATTGT
AATCATGTCTTCACTAGTGTTTCTTTCTAGCATGATAGATTTTTGTATTGCCAGTGACCCCAACAAATTTCCCAAGGTC
CTAGCCCTGTGACTGGAACACAGTAGGCATTTAATAATAGTTGACAAATGAGTTAGTATATGTAATGAAAAATCTACCT
GCCTGAGCATTAGGAAATATAAAATTAATCTAAACTTTGCCACTTTCTGTAATATCTGTGTGTGTCAGTCAGGAATGCTTT
TGGCTGCAAGAAAACCCCACTGACAGTCTTTTAAAACAAATAAGGAAGTCTAAAAGGAGTCAGATGCTGGCCTTGGTTT
AATACAATGTTAGAGGCAGCATCTCAGTGCTTCTCTAATGGTCACAAAATGACTACAGTAACTCCTTCCCTCACATCCA
CTTCCAGTTCAAGAAGCAGAATGAGAGACAGCAATAGCCACCCTGTTTCTTCATCAAGAAAGCAAAGATATCTTAGAAG
TCCCCACTGTCCTTCATTTACTTCTCATGGGCTAGAATTATCTCAGTGGCCACCCCAAGTGCAAGAGAACTAGGGAAA
TTAGTTTTTAGCATTCCCACTCTATGTTGAAAATGAGTAAGAGACAATAATTTGGGGAATAGGTGTGGGATTAGTCA
GCCAACAGTGTTTGCTACACCTTGSAGCAAAACATTTTACCTCTCTTACTCTTTGATTCTTTTCATTGAGTGAAGAATTA
GAATAAGTAATCTTTCTTCAAGAATCAATTTTCTCTCATCTATAACTCCCATATAATTCTTATTTCCAAAATGGAATTAC
CTTCTACACCAGACAGTTACTTCAAGCTTTTCCCTGAATATTTTCACATTTCCAGAAAGCACTATATTACAAAGAAGCC
CTTCTTTTCAAGAACTCTATTTGCCATAATGTTCTCCCTAACAAGGGCTGAACAGTACGTGCCCAGATTCCCTGCCTC
TGACCTCACACACCCCTCTTCTGTGTGGCGGTCTCTCAAATTGTTTAGGACAATTATCATGACTTCTCTAAATCTCCCT
TTTTCCAGGCTCAAATTCCTATGCTTCAGTAGTTTTTTCAGATAAGCCCTTCTGTCCCTTCACAACCCCTGGTCAGCCA
ACACCAGACATACTCTCTTCAATTTACTTATGATTCATTGAACAGATAATCACTGAGTTCTTGCCATGTGTGTCAGCACTGC
TCTAGGAGCTGGTGATATGGTGAGAAACAAGGAAGGGGAGGCCCTCCTCTCATGAGGCTTACAGAGGAGCAAAGGAAT
CAGATGATAAGATAAGCGTTACACACACACACACACAGAGGATCATTTTCAGAGAGTGATTCTGTGCCAAGATGAAAAT
GACAAAGGGAGATAGGAGAGTGACAAGGGATGA
AACTAGAAGACAAGCCTTCAGATGATCAGGGAGGTTTCTCTAAGACCTGAATGGGGATAATTAAC TAGCATGGAAGAGG

Fig. 9.164

AAGAGGTTTTGAAGCAGGAGGACTTTGTAAGTGAAAGGCCCTGGGTAGAATGAGCTAGACCTGAGCCAGGAAAAAAAG
ATGCTCAGAGGGGGAAAGTGTTGGAAGATGGCCTGATAATGCACATGGGGAGCCCCCACCACAAGCTCATGTAGGACA
GGACAAGAACTCAGATTGTATTCTAAATGACAGTATTTCAATACCCATCTTAAAAATGTGTCTGGACAAAACCTTAACACT
TAGCTACACATATAGCCTGACTGGAGCAGTCCAGTTTATAACCTCTCCTTGTAGTGGGCACTGTATTTCTATGGATGCA
AGCTACAGTTATATGACTTTTGTGGAAGCCATGCTATCCCCACAGTTCCACTGTGATTTCCCTATATGCTCTTTGATGCA
CACTGCTCACATGCATGAATATTACTTGCACAAAGAGAGAGAGTAGGTGGATGTGATGCCAGGAGCTATTAGAATGATT
CACATTACACTTCCACCATATGAAAAATAATGTACAGTCTAAAGCAACACTGCATCATATCATCGAGCAGCTTGGGAAA
GCCTAAATTACATATTTGAAGTTCAGTCAGTCATTGAAATTAGATAATCTGGATGGTCCTGAAAGTGTCTTTCTTGAGA
GCTTTCCTCCAAAAGCAAAATTGAATGGTAAGGAAAAAAATGAAGTACAACAAAAATGACAGCAAAAAACCTGAAAAAC
AAAAAACAAAAACAAACAAACAAAAAACCCCAAAAACCTGAGTAATATCAACCAGGCAGGAGTTGAAGGAGAGAAGCCC
AACTGTTGGTAGATAACTTGGGATGAGCCAATTAAATATTGGTCTTCTCTCTCAGAGGCCCTAATTGCATTCCTCC
AGCAGCCACAGGCCTGGGGTTTTCCCCAGGCCTGTGGCATTTCAGACATCAGAGCTGAGAACTCCAGTGACCACCAGGCT
GCATTTTAATGCTAAGCTGGGAGCATTGATACTCTCCAAGACAGCCTTGCAAGCTCCATCTTCCACTGACTCAACGTTT
AGTGGAACGTTTCAGTGTGTTTCATCTGCTGTTTCAGTCATGCAGGCAGCAGGGATCCTGAGGGCTCTCCTCCCCAATATG
GTACCAAGTTTTGATTCATCCTTTTCTCTTCCCTATCAGCAGGTATTCTAGAAGACAGTTTTATGATGACATTCACAG
CTAGCATTAATAACCAGATAAAAAATAAACAGGATGTCAATAAAGAATAATGATGTGTGTCTGTGTAAATGTATAATAA
TAATAAAAAGAATAGACATCTGTTTGTCAAGGTACCATTTAAGGCCTTGGGGCTTTAAATGAAACTGCTTTTATAAATGC
ACCAACATATGTGACTGTTTGCCTACGTCTTTCATCCATGAAAATTGCCCTGTAGACAGTTGCCTTTGAGGTGAAAAGC
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ACAATTTAAGGGTCATGGAAAAAAAGTGTTAAACGAGTTGGTAATCTGACATTTCCCTAAAAACTAGATTCAAAACCGAA
GGGAAAGTAGAGAGAGGAATCAAACTAGCCTAGTAGGATGGAAGTAGGCCTCCCTCCTTGGAGGTGTGAAAATGCCTG
GAATAATTTAGGCAAGAATGCCAACTGTTCAGTGATGACCTTGGTTTAGGGGCAGTTATAGCATGGTGGAAACACCCCTG
GACTGGAGATCCGGCATCCCTTATGGGGATATGAAAGAAATTTGGCAGGTCTCTTAACCTCTTTGACATTCCTTTCTTC
TTTGTAGGTAAGAAGCTGGTGCTGGTTCAGTGTAATTTGAACTGTGTTCTGCATGACTGTAGGGTTCTGCTTAGTGCC
ACGTGGAGGTGGGGATGCCTTGGGTGAAGATGAAGGAGGTGTAGTAAGCAGGTGGGGCTCTGCACTCTTTTTTACTCTCC
TTCTTCCAGTATCATCAGAACAACCTCCACTTTTTTGTCTTTTATACACTGAGTTATTTATAAAATCCTAGTTTTTTTAAG
TGTCAAACAATGCTTGATTATGAAATGATAGCATTATTCTAGCATTCTAACTAAAGTGTGTAAATTATAACAACCTATC
TTTAATGGTCAACAAATTGTCATCATTACTGAGCAGTGTTGTTAGTAACACCAAGTAATACAATTAATGAAATCCTTGA
AAGTAGAACCTATCAGTGAGCCATGCATACAACCTGATAAGCTGTGGATAATGTATAGAAGAATGAGTTTATTTGGTAAT
TATTTGTACTTTTCAAAATATTGGTGAACTTTCATCTTTAAGGAGCAATGGTGTCTAGTTATTCTTGCCACAACCTGCA
TGGAACCTCTCGAATCTGTTAGTATGTCCTTGGGCACAAAAACAGTATTTTTTGGAACTGCTCAAAAATTATGTAGTTT
TGTAAGACATTTAGGGAAAAATCTCATAATTTATGTTAAACACTCAGTTGTCTTGGTGTGATTGAAAGCAAATAATACG
TGCTTAACATCCTGCTAGTTGCTGAGGGAGATTTCATTCATCATTTTAGAAAATGTTTCTGGGCATCAGTCATATAATAA
ACACTTTCCAGCACTTGGCTTCACAAAGGAACAAAACAGACAATGCACATTCTGCCCCCTAGTGGAGCATATATTCTAG
ATGACAGACTGAACACCTAGACATATATGACAAGCTCTTCTTGCAAGGAGAGTCCAGTTAGGAAAATGTCAACACCTA
AAACAGTTAAAGGCTAAGATAGCTTCTGTGGACACTAAGTGCTACATAGTAGTTCCAGAGGAGGCAGAGAGTTAGACCT
GGCTAGACACTGGACAGGGGGTAGAATTAGACAGATAGTTCTTGGCTTGGTGCCTGGTACATAGTAGTTGCTCAATAAA
TATGTGTTGAATAAATGAAAAACCAGGACTGACTGAATAGACAGTTTACACATTTACATAATAAGTGTGTAATTTAGGG
AAGTGGGGTTAGAATGAGCCTGGGTGAAGGAAGCCTTGAATGAGAGATGGAGAACTGTAAACGATAAAGGTTTTGAAGT
TGGGGAAAGGTGAGGGGGAAGTGAAGTGAACATGGAGAAGCAAAAGTGTGAATATTTTCCAAGATGCCTCGACGTTAT
GATAGATGAACCTCTGAGACAGAATGTAGGGGGAAGATAGAGGAAAGAGCCAAAGAGCCTCTGAGGGTGCAAAGAAGTA
AAGAAAAACCAGGATGTATGATGAGAATTTTATTATGGAGCTATCCAGTGTGGATGTGACTTCTCTTCCCCATATCGAC
ACAAAACACTTAGCAAAACATTTCTCTTTCTCTCTCTCATGCTCTCCTGTTCTTTCTCCTCTATCTCTATCTCCTTTT
GCTCTATCTCTAGATATCTATTAGACAGATGATAGATAGACAGATACATAGATAGATAGGTAGATAGATAGATAGGTAG
ATAGATGATAGATAGATAGATAGATAGATAGATAGATAGATAGATACATCTTGGAAGTGGTAAGACAGAAAAAGTAGTG
AGTATAGCCTCAAAATGGCTGCAGGGCAAGGTAAGAAGCTGAGCTCAGCCTTCAATCCAATCCCTGTTTACACTCTGCC
TATTTATTTGTTTATAATACTCTGACTCCCCATCGACATATTTACATTCTTGGTAGAAGAGATTATTTTTTGATATCT
CTTTGTATTTATTTCTTAGCACCAAGATAATTTTAACCACTATTTTATTATTGTAAATTCATTGTTTTTACTCCAAAG
AAATACATATTTGTTGAAGAAAAATTAGAGATACAGATAAGTTAGGAAAATAATAATATCAGAAAACAGGGCCATCATT
TAAACATGGTTACATAAATAATAAGTACTGTGCTAAGTATTGTAATGATGCTGATGCTTTTGTAAATAATGATACTGA
TGGTGAAGATGGTGATGGTGATGATGATGACAATGATGAACACATACCTCCCCTGATTATGTGTAAAGGCACTGTTCTA
AACCTTTGTCTAAATTAATGCATGAAGTCTTCACAACAGCCTCAATAGATGTTATTCTTCTCCTCCTCTATTTATTTTA
TTTTAGCATATTTTATAGATGAGGTATCTGAAACAGAGATAAGCAACCTGGCCTGAGTCATACAACACAAAGTGATGGA
GGTAGGAAATGACCCAGACGGTCAGGCTCCAGTTTTTTGGGCACTACCAAATTATATTAATAATACTCTGTAGCAG
CCTGCTTTCGAGTTATGGAATAGTTTCTAGAATAAGTCACACACATAGATTTTCGAGCCCAGGACCATCTACTCTGAGCA
GAAGCTCCTACCTAACCTTGACTATAAACTGACTCTACCAGATAATCACACTTGGCCTTTCTGTAAAGTGATAAGCAACT
TGCCTCTGTGGGCACCTACCCTGAGAAAGGTAGTCACCCTGCTCCATGCTCTGTTTCAAGTACATTTATTATTATTTAT
GAATTACTTATTAACATAAATTTATTATTATTACGTATTATCCAACCTATTAAAGTGATGTATATGTATGCAGGTGTGT
ATGATTATGAAATACAAGATGAAAGAACTGTAAGAAATAGCTTTGAAAAAATTAGTAATGACTGGGCCATAGCCCAGT
GATAAGGCATGTTACATATAACATTTCTTTTCATGATGGTAACAGCCCAGCAAGGTAAGTGTATCAGGGCTGTCATG
TATTCTAGTACTGTTTGTGTACTGAATAAAGGCATTTGCCTGAGGGATGATAGGGCTCAAATCCAGGCTGCACTCCTCT

Fig. 9.165

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TGATGATCTATGAACCCAAGAAGATGAGTCTACCTGGAGGGAAAGGCTTTTCCAATATGCACAAAGTTCCACAGAAGCT
AGAGTTGTCTTGGGAATGATTATCTCCATTTTGTAAATAAGGAAATTAAAGATCAGAAAGATTGAGTAATTTCTCAAGA
ATATGTATCCAGCAAATAACAAATCAAGGACTCAAACCTAGGCCAATATGGCTCTCACTTCCTTTCTCCAAAATCACTG
CTATGTCTTTCAACTAAATTGCATTGCTCTGTTCACTGGTACAAAGCTTGTGTCACCATTATAAAATTCAGGACTATCT
GATTGTGTATTATATTTTTTAGATCCTTTCTGAAAGCAGGTTTTAATGGGGAGGGAGAAAGTAGCAACTGGAAGAGAGT
TTTGAGATGGTAGATGGTAAAATGTGCAAGCAAATTCCTCAAAAATCTGGGGACACAGACTTCACTGCCTTCAGAATAA
TTGAAAGGAGAACCCTTCAAGTATGAAGCTTCAGCTGAAATTTCAACGTGGTTTATTTACCTAATATGCTGTTCAATTAA
TTACAAGCTATTAGTGACTATTAGCCTATGAAATTGTATTTACAGTTCAGAAATGTATTTGATTTTCACAAGCCTTACT
GGGGAACAATGCTCAGTCAACACTTTCTTCGCCTAAAGAGTACGATCAAAGAGCATGAAGCGTAGTAAATGTTTAGCCT
TTGAATTTGGAGTTTACAAAGGATAATTATCCACGGGTGGTGAATTCATGTTCTCTCTACTCCACACAGCCTCTGGT
TTGCATTTACAGTTTTCACAAATCAGATGTAAAATTAATAATATAATCATTCCATTTTAAATGGTTCCCTGTACAATTAT
CTCATTTTAAATCTTTTGCTAATGTGGTTATTCTAATGCAACTATGATCATTATTTTATGTATGTAAAGATGTCAATAC
TGTCCAGCTGATGATATTATCTAATAGTATGTAGCAAAGCATGTTCCCTTCCTACAGTGCCTCACTCTTTCGATGGGTA
AGATGGGAGGTGGGAGTGGGGGGAGCATGGCATTAGCTGTGGATAATAAAGCAAAGTAAGTACAATAGGTGTTCAATAA
AGTCATCATTTTTGGTACTTACGATGTAGTTTTCTTGTTCCCTTTGGTTCCTTTTGCAGTGGGTGTATTTGCTTAGACAG
AGAGAAGTGGCAGAGGGAGACTGCATTTGTATTTGGTTATTTTCAGAGAAATGCAACTTGGTATTATGAGCCTTTAATTC
TGTAGGGCAGTCTCCAGATCATTTTGATGCTGAACCTTTGTCATAAAAACTATTGATTTTTTTCTTCCAATTATAACA
TTTGCTGGTTTCTTGTCAGTTAATAACTGACATAACATTATTTTCAGACCTTTCACTTAGGGCTCTGTTGTGTGTTTCTC
TTGATCAAAATTATTGCCTTAGCATATCAACTGAATACACAAGAGAAAAATCAAGTTATCTATGTTTATCATATGCATT
TTTGAATACCTATATGTGATGCTTGAGCATAGTGGTTTTWATAATCCTGTAAGCGATTTTTTCAGGAGACCAGATCTGAC
CTCCTCTCTCACACCCCTTTTCACACTGCCTGCAACACACCACACATACCACACACATCACCACCCTGACCGACCAATGG
AATCTGAGCCAACTATAAATCTCAAACCTTGTTGATATTGGATAATGCCTGGAGGAGAAACACGGCTACCGTTTCAGTAAT
TCCAGAAAGAATTTCTTCTTTAAACTCAGTAAACCCAATGGCTTCTCAGAAAGTTCTTTCTGAGTTTTCTTATTTGAAA
TTTGAAAATTATTTGAAATTGACTTAACCTGACATCACCTGTTCCCTCTGGATCTCCAGTGATTTACTGTTGACTCTT
TCAAATCCACTCATCTCAGGGCTGGGGGACTGAGTCACAGTCTGCTTTTTCTCCTCATTTGTCATTTCCAAATTTTTCTTC
CCTCTCCTTGCTGGGGGCTGGTGCTAAGTTGACCCCATTTTTTCCCTCCTTGCTGCTGTCACCTCACTGCCTTTCCAGAATC
TTCTGCATCAGGTCAGTCACTCTTTCCAACCTAATTCCATATGTCAACCCATTCATCACCGGAGCCACGTGGTTTCTCT
CACACAAATTGTCTTCTCTGCACCACAGATGCCAGTACACTCTGGCCCTTGCCATCTGTCTTAAAGATGCAACACCGGT
CTGCTTCCAAATCCCTAGTTCAGACAGCCTGCCTTCCAGTTGCCACATCCTGTTTGTGTTCCAGTTTAACTATGTAC
TGCCACTGCTGCTTCTACTTGTGACACCTTCACTGACCTCTCCATTTCCCCCATCCCTCCATTTTCATCCTTACGTTCT
TCCGGCCCCCATCTCTGCTTCAAYGAATGAACCCGTTTTTCATCTGTTTTTCCATGTTGGGATTCTAGTCAGATTTCTCT
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TGAGATGATCTTTGTCCAATATTGATTAGTGGTGGAGTCCCTTGATAGCCAATCCATATTTCTTTACAGTAATAGAA
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AATACTCTAATTTTAGGAATCATTGTTACAATGTTTCAAATCTACTGACTTACACATATATATCCTACATATATTGTCT
AAACATATTCTGTAGAAGCAATGGGGATTGATGATTATGAAAAAAGTGGCCATTGCTGGTTTAGGCAGTGTGTAAATGT
TGCTACTGACATCCTAATGTGATTTTAATTCTTGCACTGCTGGTGTCATAAAATAGCTTTCAAAAAGAGCATTTTGATC
ACGACTCATTTTCCAATTTCTCTCCTCCATGATCCCAACATCTTCTTAAGAAATCCACTCTGTACCTGAGTTTCCACAT
GGAACCTTAGGACAATGATGTTAGTGAATTAGAGGTCTGGCAGAATCAAACGAAAGTAATCCTCACCAAGTCACTAAACC
ACACTACTTTGAGTCTATGCAAGTTCAGGAATTTCTATTTGAGCAAAGCCACAAATGGCCAGAGCGGACCTCAGGCTT
TGTGCTTGAACCTGCTAATGCCATTTGATTTTTTCGATCCCGCACTTTTCTTGCCAGACCAATGGTCTTTTTCTAGCAGT
AAAATTTCAGAATAGGTAGGCTACCATCACTCTTTGGGTTACCCAGTGTTTACTTAAGTTGAAATTCAAATAGATATG
TATTTCTATATGAATGGCTGCCATTCTGCTATGCCCATGTTTCTGTTTCTCACTCATCTCTGGCTTGTCTCGTAGAGCC
CTTTCTGGTGCTGACTTATGTTGCTGAGTCAAAGGCTTTCATAGTCAAAGCTCTTCTTTTTATCTGATATGTTTTGCAG
TTTGCTGTACTGCGTCAACTCTACTGGGATCATATTTCTCTTGAGACTAACTACTAGGCAACCTGTGCCTTCTTCTCTGC
ACTGCATTTTGCAAGTGCTTCTTAGGATTTCACTTTTCAATTAAAGGCAGTAGATAGCTCAGAGAATGTGGAACAGTAAC
TAGAACAGATCCAAGAGCTGTAGAATCAAAGAAGAGTTAACTACTGTCTCTGAACTTAGAATGTAACCAAAGTCAAGC
TATGCAACCTCACTAAGCCTGTTTTCCCATATTTAAAGTGGATTGTAATGTTTATGAGTCTACCACTACATYAGATTG
GTGCACTGATTACAGGAGACAGCGCATGTAAACTCCTGGCATGGAGCCTGATGCATTTGAAACATTTACTACAAACTA
CCATTGTTGTTATCATTATTAATAAATAACCCATTCTTTCAAAGAATCTTGTTCTTGGCCACCAAGCTTGGAAGTCAGTG
ATATTGTTTCATCTTATTTGCCCGGTTTTTTTTTTTTTTTTTTTTTTTGGAGAATAATTTATGACCTCTGTGGCAGCAAG
TCTGGCCTCATCTTGACCATTCTATCTGCATGTACTAACTTTTCAATTTTAATGGCCCTGTGAGCTTTCTCCCATAG
CCAAAAGTGCTGTTTTTCTATTCTAGTGTTTTCTTGTCAGGCAACAATAATTTCTAAATAACATGCCTCTTCTAAACC
AATCTCTTAAAAAATAGATTGTTTTTGTCTGACCTTTCTCATTTCCATTTTCAGAACTATTTTTTCTCCTTCTTTAA
GCAGGTTACAAACCCTTATATTTTTTAATATGCTGAGGCTTGTAGTCAAACCATTAAGTGTCCCAAAAGAAAGCCCTTTT
AGCACATTTACAGCTTCTTGCTATCTGTGCCCTAACTAAATTGTGCTACAGGGATAGTTTCTTAACCATTAATAATGATA
GGTGAACACAATTTTATTTTGAAGGATTACTAAGCTATTTTTTAAATTTCTATAGTTAGAAAAGAAAACTGTTTAACT
GAACTTGAACCCAATTAAAAATAACAATTCTCATGGCTCTCCGTTTTGCAATACTTTCTTATTCATAAAAATAAGAATGT
GCACTTTACCTTTTTGACCCATATATTTGATGTTTTTCAATTCTCTACACGTTCTTTATCTCTCTGTGAGTGTGGAATA
TTCTGATTTGTTAATTAATACCCAAAGCAGCAGTTGGCTATTAATGAAATGTACAGAAAGGTATTTGGTAAAGGCTACA
TGAAGATGATGATGACAATAACAAAGATATCTGATTGAAAGTAATCTTTACCTGTTAATCTTCCCTTAATGTCTCCGAC

Fig. 9.166

Fig. 9.167

Fig. 9.168

GTTCTTGGCATATTTTATGAACTCCTATAAAACCCCAACCAGGTTGTGAGTAGTTATCGCTTACTGAGACCTTAAACACA
AACAAATATTTTAATGTTTCATTTATTATCCTGAATAAGATACATAGATATATATTGGGAGTTGAGCGCTACCATGTCAAG
AGGAGCAAGCCCATAGAAATGATTTTTCTTTTCATTTTTTTTGCAAAACAACCTGCAATGTCCCATTAAGACTTTTCCTGCAT
GTTTCGTGTCAGATGAAGGAAAGTATTTTCCACTGATACCCCAAGGTTGTTAGCTAGTCCCACGACTGCTCTTCAAGAAAC
ACAAGAGACCTATCTGAATGTGGGTAAAAGGATTTTCATGTTGCTTTTTACTTAGAAACCCAGATTGTTTGTAATAAGCT
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AAATGATAAAAGCATCTGAAGTTCAGCAAAAGTGAGCCTCATCTGAGCTGACTTGTCTGATTGTTGGGATAGCCTGAGG
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Fig. 9.169

ACTGCCAGGTCTTTTCAGTCTTTATTTGAATAGGTTTCAGTAAGAAAGAATTCACCTCGTGCTTTAGATATGTATGGCTTTT
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Fig. 9.171

Fig. 9.172

GA CTCTGTGTCTATTCCACTGGTTTGGAAAATAAAACAGATTGATTTTCAGTCATATGAAGATACTTGAAAATAGGTTAC
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TCCATACCACAGAGGCTCAAGGTAGA ACTGAGCCCCCTTCCTTAGTTTTTCCCAACCACAGCCTTCCCTCTGCCCCAACA
AAACAAAACCTTGTTACACATGGGATTTCTGCATTGTACCACACTTTACTGATGGTTTGAAGAAATGGAAGCAATGTTT
ACCATAATGTGAGAAGTGACTGTGTTTAGACTTT CATATTTCTTAAAAAGTCAGTTACCCAGTGATTTCTATATGGAAGG
TGTTAGCCTTTTGCTAGGTTTGCTTGGGTTTTTTCTTTTCACTTCTACCTTCCGTACCTCTCCTACTTCAAAAATTGCT
TATAGGAACTCCAGATTTTTCTTACTTAATGGCATTTAGTGCTTTACGTCTCTCCTGCTCCATCCACCTTTTAACTCCC
AGAACTACGCAAGTGCTCAGCAGAACAAAATGGGCTTTCATGTAATATTACCACA ACTTTGATAGAAGATATTTTTTGGTT
ATAAATAGCTTTTTTAAAAAAATTTTTTGGTGATTACATGTGGGTACGATGACTAAATTGAATTTTGCTCTTCGTCTCAT
ACACACAAACACACACACACACACACACACACACACACACACACAGGCATTTATATATAGAGAGAGAGAGATG
GAGACTTGTTCTGT CATCCCAGCTGGAGTG TAGTGGTGCAATCACAGCTCACTGCAGCCTCAACCTCCCAGGTGCAAGT
GATCCTCCTGCTTCAGCCTCCCATGTAGCTGGGATTACAGGTGCGGGCCACCACACCTGGCTATTTTTTTATTATTTTTT
GTAGATGCGGAATCTCCCTATGTTGCCCAGGCTAGTAACTCCTAGGTGTAAGTGATCCTCCAGACTCGGACTCCCAAAG
TGCTGGGATTACAAGTGCAAGCCCCCATACCCCCAGCCTTCTGCTCCTATTTGACCTAGAAATTC CATATAGTAGCCAT
AGCATTCAATTCACTCAACAAATAATTATTGAACACCTACTTGCAAAAAGGAATTCAGTTCCCTATTCTGTTGGGGTGATA
ATCTAGTAAATAATAATAATAATAATGAAAAGTGTCATGCACCTCTCTAGGCTCCAATAATCCCTATGAAGAGAGGGT
ATGCTTAAACAACAGATGCTTATTTTTCTCACGGTTGTGGAGGCTGGCAGTGTGAGATCGGGGTGCCTGCATAGTGGAGT
TCTAGTGAGGGCTGGCTTCCCTATGCCTTCACCCAAAGTTTATGTGTTATTAATATCAGCAGCCCTCTATCTCTGGAT
GTCTCTGGTGCCTTAGAAGACATATTCTCTACAGTTAAAGAGTGATCCTCAAGAACGGACAGGGTAAATATATTTTTATA
TTGCTAAATGCCCCCATTTGAAGGGCTATATACCATTTCATGTAGCAAGGAGCACTTAATGGCTCCAGCAGAACAAAAT
AACTCTACAGAAAATTTTTATTGTGCCCAAAGCATCTAACGAGCTGGTCCCTCCATGCATGTGGTGGCACTCTTGGAAC
TGCTTCCTGAGATCCTCTTAGTGCAAAGCCAACAACATT CAGTACATTTCTTCACTTGCTTTTGGCATGTCCATGAC
AATATTTCCAGCTGATCTAACTTCTAGAAAATCTATTTCCCAAGCAACAGGTCTTCTATTTTCCCTGCTTAAGAGTGT
TTATTTTTTGCTTTTTTTCAGCAGCTTCACA ACTTCTGACTTGGTTGACTTTTTTTTTTTTTTTTTTTTAGAGAGAATGTCT
CTCCTCCATCTGCGAAGTGCTGCATTTTAGGTTTGGTCTGACAATTTGATGCAACGGCTCAGGTATATATGACAAGGAA
ATGGAATTTCTATCCTCGAAATCAATTCTGAACAATAAAAGCTGAGAGGATAAGTTTATTCAAAGGTGATTTGGTATTG
CGACAGGCTGCTCCAAATCAGAACTCAGCAACCTGATTTTGAAAAATTTCCAAGTTCATGAAAAGAATCACTTTGTACA
TAAGGGCTTTCTACTTTGTTTGGGTTTGAGCATAACTAGATAGCTTAGTGGGATATCCATATTTCTTGCA TTTGATGATCA

Fig. 9.174

TATACTAGACTTTGAAAATACTTTAGTTGCAGTGCATTATTTTTGTGTAGTCAGAGTACCCAGGGCAGTTTCTCACACA
TTCTCTCGTTTTGTCTCATAACATCCCAGGAGCTGGATATAATCTGTGATTTATGAGGAGAGAAAGTGAGAGACAGAGG
TGTAAGTGAGTTGCCTAAGGTCACTCAGAGAGTGAGTGGCAGAAGCAGCACTTGAATCAAGTCTTAGGATTCTAAGCCC
GGGGCTGTCTGCTCCCTCCTAGACCTGTAACTTTAGAAGCAGGAATTTATTTTCATATTGCTTATATTTTCAGTATTTATT
TTTAACCCCTTACAGGGCCCCGGTCTTGTGACTGACTGTAATTGAGAAGGTACTCTACAAATATTTTTGAAAGAATAAATA
AAATGAATACTACAGACCATGTTAATAATGTCAAATAACTCTGGGAGTTTGAGGGATAATTCTGTCAATACGAGTTGAG
GGTGTGCTTGGCCTCTCCATCATTTACCCTGCTTTTATGAGGGTTAGCAGCACAATGTCTTCATTCTAAAACACAGAAT
CATGAGGTTAAAGAAAGCATGCAATGCCATCAGCTGAGAAGAACCAGAATTCACAGTGCAGGTTCTCTAAAGTGTTT
GTGTTGCTCTCAGCGCCGTGAACCTCCACCATTTATATTCATAGCTTTTTTAGCCTTTGGCTAAGATCAAGTAGAGTGTTT
GTTCTCCTGGCTTTTAATAAGTAATGTGTTCTCCATCCAAGGACAACATATTTAGGCTGTACATGCATTACGAAGTATT
AAGCTCCTGTAACAGTAAATTTCTCAACAGGAGATTTATGTGAATTTCTTGGATGTATTCTGTGTGACATTAGGAACA
ATTAGGAATATAAAATGCCATGGCAGATTTTTCTGTGATGCTGGACGTACTGTCAGGACTCTGAGTCTGTGTACTCGGA
AGTCTTCCCTCAGACAGGGACAAGGCGCATTTCTTTTCTTAGGAAGAGAATTAACAGTGCACCTCCACCTCCCTGTGCTC
ACCACATGGGCACACATCCCAGGGCGAGGGTGGAGTGAGCTCAGAAGAACCCACGCTGAGCACAATGAAAGCAAAATTA
TTATCAGAGAAAGAAACAGATTGCAGCAACCTGGGTTTAACAGGACACATAGTAGTAGACAGGAAGTTTTTCGTGATCT
ATAACACATTTTCATATTAGGGTCATGTTAATTTCACAATGAATTGCATATTAGGCTCAATTAGAGAAACAATAAGAA
GGAAGACGGGATGACAGAAAGCATTGAAGAGGAGGAAGCAGAATGGAGGCAGAGGGTGGAATCAAAAATAAGGACAAAG
AGAGGAAGGGGAAAGAGAAAGAAAATGAGAAGTAACGTCAGCAGCTCCGTAGGTTCTGAGGTGAGTTAGCTGGTCGTAG
GTCAACAGCTTTAGTCACACATCCTGTTATTTCACTTTGGGCATCAGAGCACATTTTCACTACCCTTTGAAATTTCCATA
CAGTCTACAGCGATTTTGTAGCAATCTTTCTTTACCGGGAATTTGCTTATCTGCTTTTCTTTTCAATCTTTGAGGTCAA
CTTATTAAAGGCTTTAGTTTTCACATGAAGCCCGAAATATTTTGAAGGTAAGAAAAAATGTGCCTTTGGTCTTAAATAT
TTCCACTTAAATCTTGATTCTGGGCTACTTTAATAATTCAAAATAAGAGTGAGTGGCTTCTTTTTTCAATAACTGAAA
TACAACATTTAAAAACACTCATGTAATTATTTTGTGTTTCAAGTTGTTTGAAAGTAGACAAGGTTTGATTGTACAGAG
CTGTGTTTTAACCTGGGCTCTACATGTCATAAATACTGTCCCTGTAAAGCTCCTGTTTCTCCTCTACAGATAGAGG
TACTCCAGTAGCTCTCCAGGCTCTGTGTGTCTCTGTACAGGTTTGGCATAAAGTAGACAGGAATGGTGTGAGTTCTC
CCTCCCCAACTCTCTCCAATGACACTCCACCAAGTGACAGAAACGTACAGAAAGAACTCTTCTGCTTTTCAACTTCC
ATTGATCACACTGAAAGAAATATCACATTAGCTTATAAGGCTCAAATTTATCTTCAGTGCTATAGCTCTGTGTACTCTA
AAATCAGAGCAACATAACTTTGGTTCTGGATGAAATCGAATCAGGACCTGACTCTACTCCTAGAAGAACGCTGACCCCA
AACTCCTGTTTTTGCGAACCCTGAAGGAGCAAGTAAAGATGCCAGCCATTAAATATGAAGACCTTCAGAGACCTAGAAC
TAATGGAATAAATCAAGGATTTCTATTTTATCTTAGATTAAATGCCATTTATATGCATTCAGGGCATAAGTTTTCACT
CCCATGAGGAATAGATTTCATACCTGTATAAAGCTGGGAATATAATGACTAATTAATATACTGTATGACTTCAATATAG
CCAAGAAAATTACAATCATTTCCAAGTAATACTGTTTTTCCCCAGACACAAATCTGAGGATCTTGAATCTTAGCACTGGA
AGGAATGTAGACATCACGCAGGCTCATCACTTCTCTGGCACAGAACTTGACTTGCCTTGAGCACATGTCTGGTAGCTTC
TTCTGGGAACATAGATGACATAACGCATATTGCCTTTCCAGGCAGCTTGATCTGTGATGAGACAGCTCTGGCCTTTTA
GAAACCAGCCTTTATACTGAGCTGTGAGCCTGCTTCTCTGTATTGCACATTTTTTGTCTTAAATTTCTGAACAAAAAGTT
TTTGTGAATTTTCTTCTGGAATGGTCTTCACTTGAACAACCTGTCATGCCCATCTGAGCATATTTCTTAGGTGA
GGCTTCCCAGTTTGGGAGACGCTTCTCTCATAAAATATTTCTGAATTCATCAGTATCTCTGTTTCTATAAAACGCGTGG
CTGTCTACTTCAGGCAAATCCTAACCAAGCTCAGGTAACAGTGAAGTGATTGTTTGTGTAACAGTATTGTACTGTT
GAAGTTAAAGAGGTCTATGATTACATTGTGTTTTATTTTGGATTTTGCATTATTGAGTATTTTTTACTTTATTTTCTTTT
TGTAAGCCCATATCACTACCTATAACAAGCTTTTGATAAGTGAACCTTTCTAATTTTTCTTATTTCTATAGCTATGTAAG
GGAATCTTTAAATTTTGGCAGGACTTTCTAGCCTAATGCAACAAGGGAGGTTACATTAACTATAATAATTTAAATTTCA
TTAGTTTTTTCCACATTTGCTGAAGCAATTTCTTCTACACAATCTATTTTCATTATTTCTGGGAGTTCTAAATGTGTCCC
ACAGGAACACATCCCCCTTCTTGAACCACTGCTCTTCCACCAGAACCAGCAGCTCTCTGGACTCCCCAAGGCTTATTA
GGTCTCATTTGTTACCCAGGGAGTCAAGCAGCTATCACAAGGCCATGCTTTGGCTTTGTAGCTGCTCCAATGGATGTTAA
AATTTCTCACTTTGTTCTCTGATACTGTACGGAATTATATGCAACATCTCTTTACTTTAAAGCTCTCTGCTTCTCCCAGG
ATCCAGAGATTTCATCCCTGGAAAAGCATTCAAGTGACAGCTGCCATATTTCCATTTAGAATGCCTCCTTCTTCCAGACAGC
CAAGGTTTTGTTATTGGGCTCAGCTCTATGTGCCAGCTCCTTCTTCAAGTACAAGGAGCTCTCTCCAAAGAGGTCACCTA
AATCCCCTGCAAATAGTCTTTAACTAGACCTTTTCAAACCTTGTCTCTTTTTTTTTCTTTGAAATGACACTGTACCATGA
ACATTTTTATGAAATCATTAACATGTTAACTTCATCAAAGCCATCATTTTAAAGTCTTCCATGATGTCTATCTTATA
GAGATACACCATACTAGAATGCATTTTTTTTCATTTTTCTATTGCTGAGAATGTTGGGTTTTATACATAGCTTGTATTAT
TAAGTAAATGTTGTGCTTTACCTTGTTCTTATTATACTTTTAATTCAGCCTACTGTTAGCATCTGTGGATGATTTTCTA
TGCTTATATGTCACTCAATATATTAGCATCATTTCCAGCTTTTGGCTTTACATATGCATTTTTTAAGATGTGTCTCTTAC
CTTTTAACCATCAAATCAGATTTTAGTTGCTTGATAGACCAGCTTTTATATTTTCAGTTTATGCCACCTCTAAGGTCGT
TGTAAGTGAGATGCCTATATAAAAGACATACAAATTCAGGAGTGAATTAATAATCTCACTTGAAATTTTAAATGTCACTG
AATATTAATTGTGGTCGCTAATTGGTATATTTTCTAGTAATCACTTCAAAGCTCCTCATTAATAAATAAATCATAAAAA
TATAGAATTATAAGAGCTTTTAGGTATAGTTCAATGATCTAATTCATCCATTTATCTGTCCATCTATCCTTCCAGCAAG
TCATTTATCACCTTCTTCTGTATATTACAGATGATAATAAAGACATCTGGGACAAAAAGGTAAGTCACAGACCTACCC
ATGTGTATGGGTGATAGAGTAGTGAATACAAGTGACAAAGGGTGTGATTCTTCTGTTTGTAGAGGTGAAGAAAGTATTC
CCAAAGCAAGTGACATTGAATCAGCACCTTAAGATAAAACGAAGAACTCAGCAGAAAGAAAGTGAATGGGAAGATGTGG
TAATAGGTTTTAAATGGACATGGCATGTGCGCAACTAAATTTCTGTAGCTCCAGAATATTTCTCCCCAGTCCCCGTGAGACG
AAGTTACAGAGCTGTGAATCCGGTAGTGAAAGCCTTGGTATAAGGCGCATAAGAAATGTGCAGATTATCTACCATGAAA

Fig. 9.175

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TAGAGGAAGAAATCAAGGTTATTCAAGTAGAGGAATGACACGTTTTAGAAAGACTTTTGTATAACGTTATGGGAAATG
GATTATAGGCTGGAGGCAAGAGATAAGACATTGTCAGTGTGACAGAATATTTTATTAATAAGCACATAATTTATACTC
AGATCTATGTGAGATGAGACCAATTCATGTCAACATGTCCAATGTTTTTGAAAAAATGCTAAGTTTTTATTATCGCTTT
TGTTTATTTCCCTGTGAAATATAGTAGGCTAGCCCTCTATTTCTCCTGGCGAAATCTTTAGATAGGTGGAAAAAAT
GTTGAAAAAGGGCATCAATGACCTCTTTTATCTTGGTTTTATCTGCCTTTTGGAAGATACTTGATTCAAAGCTATGAAA
AAGCATCTCACCAATCCCGTATTTCACTTGGTTGGACGATGTGTCAAGAGAACGCCTGAGGCTTCCTCTGTCCGTTAAC
AGGCCAGTCATGAGTCACAGTCTGAACATGAGGAAATGTTTATGTACTTTTTTCATAGTCATGGCAATGAGACTTTTTTG
TGCCCAAAGTACTATCCTCGCCTAAAGTCCCTGTAAAAAGGTTATTTTCTGCCACAGGCAGGTAGCAGAATGTCAAGA
TCCAATTCCAAGTGCTTCATCATCACTCGGATAGCCTCAGGAAGCTGAGGTTTTACAGCTGCCTGGAACCCACATGCTT
CATGTGATGTAGAGCAGCCCTGTTAATTAGCTGTTTCAGGATTCTTAATGCCCTATGGCTTTGGAAACACTTGTTAGCC
CACAGAGGGAGGGGAAGAGCTGGAAGCAATTGGACTACAGCCTGAATTTATTTTAATTCAATTCAGTAACCACCTTATC
TTTCTAAATCCCTGTTTAGAATCTGTTGAAAATTATAGTTGATTTAAGTTTTTCTCAGTGATGCATCCTGAGAGAAGAG
GGAGATTACAATCAAGGTTTTCTGAAGCCACAGAAATGTGGAGTGCTGCAGAAAGAGCTGATCGATGGAACCTTCCAG
ACACATGTATGCAATTTTCTAAAGGCTAGAAAATAGATGCCGGGGATGGGGAGAGGGGAGGCCAGTTGATGTGAAAGG
CTTGTGTAGAGTGCCACGAGTCTGCATTTCTGTTACCTACCAGAAGAAACACTGCCAGTTCCCCCACCCTCACCCC
CCAACCATTTCTCACTGCAGAGAGCTGGCACTCAACAACTCATAATTTTTCTCAATACAATTCTCATGACATAACCTGA
CCAACCATATGAAACCTAATGAAGAAAACAACCATTGCCATTGGCAACAAATGCAGGAGTTGCATCACAAGTAAACCAC
ATACTTTACTTCACAAATTTGAACATTAGAGTACCACCGGCAAAAACATGGTACCATCCTTCGGTATGGATGAGATATC
AGTTTACAAAACCTTCACTGGAAAGTCCATACCAAAATTTTTAAATGCATGCAAAAGCTAATTGTGTTGACATCTTTC
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AGCATCCTCTTCCTCATATTCATCCTCTGAAATACCAAGAGGTGAAAGAGTGTTTCAGTAGCTCACCATTTCTGGCTGT
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TAATACATTTCTTAGAACCTAGTAGGTGCTCAATAAACGTTAGCTGTATGAATGAGTTAGTGAATCAATGAATGAAGTGT
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CAAGACTGGGTAATTTATAAAGAAAAACAGGTTTAAAGGACTTACAGTTCCACGTGACTGGGGAGGCCTCACAAATCATG
GTGGAAGGCAAAAGGCATGTCTTACATGGTGGCAGACATGAGAGAAAATGAGAGAAAACCATGAAAAAGGGGTTTTCC
CTTATAAAGCCATCAGATCTTGTGAGACTTATTCATACCATGAGAACAGTATGGGGAAAATTGCCCCCATGATTCAAT
TATCTCCCACCAGGTCCCTCCCACAACACATGGGAATCATGAGAGCTACAGTTCAAGATGAGATTTGGGTGGGGACACA
GCCAAACCACATCACCCATTTTAAAACAGGCTTGATATCAATTTATTGCTAGAAAACCTATAATTTGTATTTTCTTTTAC
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CCCAAACATGTTTTCAATACTAATATTAACAAATTAGCTTAACAAATACAAAAGGTAATGCCTGGTTTTATACAATAC
GGATTTGAGAAATGCAGGTTTCAACCTGTTCCACACCATGCCCTTCATTATATTCTTCTGCCAGAGATTTTATTCTTTA
CTTTTCATTCAATCAGAAAGAATTGAGAGGAATTTGAACCCATTCTGCCTATTTAGAATATCATTTGTACTGACTTTAA
TCTTTTGCTCCTAATCCATGAATTCCTGAGGTCTTTTATTTACATCCATAAATTTGAAAAGCACAATCCCCTTTTTTTT
TTAAAAAAGGAAGTTACTTTTGGGAATCAGAGACATGTACCTCTTTGTTCTCTTGGAGAGCTGTGGGTTAT
GGGAGGGAACCTCATCCTGTTTCCCTGATATGCAGTGACTTCTCTTACACAGATGAGTCCTAAAACCTTTGTGAGC
TCAGACAAGTTGGCTGGTTTTATTTTTATTTTTTTTATTTTTGAGATTGTCTCACTCTGTCACTCAGGATGGAGTGCAGT
GGCCCAATTACCAACTCACTGCAGCCTTGACCTTCCAGGCTGAAACCATCTTCCCACCTCAGCCTCCCAAATAGCTAAG
ACCACAGGTGCACACCACCATACCCACTAATTTTTTTGTAGAGATAGGGTTTCGCCATATTGCCCAGGCTGGTCTTGAAC
TCCTGGTCTCAACCAATTCACCTGCCTTGGCCTCCCAAAGTGCTGGGGTTACAGGTGTGAGCCACCATGCCTGGCCAAG
TTGCCTGGTTCTTTACATGGAATGTTCTTTACATGGAATGTTCCAGGATCAATTAAACACAGATAAATAACAACCTTATG
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ATGTTTTAGACTAGGACATCCCAAGTGAATTGAAGATTGGTACAGGAGCCTGAGGAAGGAAGTTTAAGAGAGTGAAAGC
ATGAGAGAAGAGCCAAGATTATGTAGCCAGAGAAGGAAGAATTATTACCTACCATTACTTTAAGGGACTGTGCTTCC
TCACAGTAGTGGAAGACAGATGCTCATTTTTTTATTCAAGACTGTGTAAAATAAATGGCCTTAATTTATAGATGTGGATG
CTGGTTTTATATCAATACACCTCAACCTGGGATGGCTCCAAATCTTGACTACACATGGGAATAACTTGAGGAGTTTTACA
AAAAAATGATGCCTGGGTACACCCCTGAGAGATTCTTATTTATCTGGTCTGGGGTGGTGTGGCCAGGCACTGAAATT
TATAAAATTTCTCCGGGTGATTCTAATATGCAGCCAAGGTTAAGAATGCAGCTGTACAGCTGTAGATGGAAGAATACCAA
AACCAGGCCTTCTGCTAGTGCCTGAGCTTCTCCTCATTTTATGTTTCTGTGATGTGTTTCAACATTTGTTTCTAGAATCTC
CTGGGTAATATGAGTTATAGTCTTTTGGATGGAGTAATACATTAACACATCCATACTCACATAGTTTGAAGGGGCCAG
GATGCAAAGGGAAGTTGGAGGAGGAAGAGAAGGAATGGTAAAGTCCAAAGCATGGGTGAAGGGGGGCAATATAAACC
CACAGAGCAAAAAGGAGATGTTGGGTCACTCATATTCTATTACATAAATTGGATTTACATTTCTTGTCTTTTCAATTGCTG
GTCTTTATTTCCACAAGTCTCCAAATACAGTCATAAAATTTCTCATGAGTTGTTATAGCAAAATTGCTTATCATTTACTA
TTTCTTAAATGAATGCATAAACTGTATTACTTTGGCAGAAAGGATGCTGCTGGGTATCATATGTAATGTATACCTAGTAA
GGTGGACAGGACCATTAGGACTTTAAATTCCTTTATAATCTCAAAAGTCTGTGATTCTGATCTTCTTGGTCTTGCAACT
CCAGAAAAGGTGGTAATCACTGGAGTAGGCTATTTATGGGCCTGTGAGATAGTGAAACATGCTATTAAGACAAAATGAG
AGACTTCCTTCTCTGAAATGGTTCATATAAAAGTAATATATGGCTTAAGTATCCAGGGGCTCATGTAATCTCTCCATG
TCAGTATCTTTTACGGGGAATTTATTTAACCAACATGTATTGAACACTTACGTGCTACACTAAAATCCCAGATTTGTCA

Fig. 9.176

AGTTCCCGTTTAAAAAGTGGGTACTAAGCCTAAGATATTAGCAATACTTGTATCTGAAAAGGCCTCACATCTAGAACT
ATAAAGAAGCTTCTAAAAATCAATCAAAACAACATAGTTCTTAGAAAATCACAAAGGAACAGGTTTTTCAGACTCGAGGA
TGTCTATATGGTGAAAAAGCAGCAGTATTTTAGAGTTGGTCTGAACTGACTCACAGGAACCTATTTTTAAATGTCCAGA
AATTTGCTAAGCAAATTGTTAATGCATCAATTATCTAAAGTTTAATTATGTAACTTCAAGTTATCTGAAAAACATAAC
ACATACTCAAACTCTCATCCCTTCCTAAGTGTTTTACTACATTCTGTTATTTCTGTTTGTGAGGTCATTTGTTTATTG
CATCTGTATGATGGAAACACTGTATAAAATGACATGTTATTCCCAACTCCACATAGGCTGATATAACGTTGGTAGCTTA
AAATCAGCTTTAGTCCAGGTATTTATACAATGGAAATGGAAATCAGCATACGCCACAAATCAGGAGTTGATTTTTTTTT
TTTTTTTTTTTTTTTTTTTTTTTTTTTTGAGATGGAGTGTCGCTCTGTGCCCCAGGCTGGAGTGCAGTGGTGGCATCTCGGCTCA
CTGCAACCTCCGCCTCCTGGATTCAAGCGATTCTCTGCCTCGGCCTCCTGAGTAGCTGGGATTACAGTTGCCCGCCACC
ATGCCTGGCTAATATTTGCATTTTTTAGTAGAGATGAGGTTTCATCATCTTGGCCAGGCTGGTTTTGAATTCCTGACCTC
GTAATCCACCTGCCTTGGCCTCCCAAAGTGCTGGGATTACAAGCATGAGCCACTGCGCTCGGCCAGGAATTGATTTTT
TGTTTAATGGTGTGACTGTCTACAACTGATTAAGAAAATGTTATTAAAGTTGTACATCATGTCTGCAACCATTAAAT
TGTAAGTAGCACAAAAAATCTGGGGGAAACAGTCTTCTAGACTTTCCAGATGCAGCAAAGAAATTGCCTTGTTCCAACA
TACATTTTTATTGTTTCACTTTTGTCTTACTTAACCACTAAGTAAGACATAGTTCCCTACATAAACCAATCATGTAGGAA
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TTTTTTTTTGAGAAGGAGTTTAACTCTTGTCAACCAGGCTGGAGTGCAGTGGTGTGACCTCCGCTCACCATAACCTCTGC
CTCCCAGGTTCAAGCGATTCTCCTGCCTCAGCATCCTGAGTAGCTGGGATTACAGGCACCTGCCAGCACATGTGGCTAA
TTTTGCATTTTTTAGTAGAGACAGGGTTTCACCACGTTAGTCAGGCTGGTCTCGAACTCCTGACCTCAGGTGATCCACCC
TCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACTGCGCTCAGCTCACCATCGCTTTTAGATAAGGAACT
GAGCCCTAGAGAGTGGTTGGCTCGCCTCAGGCTCCAGGACAAATATGACTTAATCAAACTATACTCCTGTTCTTTTCAT
TCACATAAACTACTTATCTAAGGATGCTGCAGCAACACTGCTGTGAGGCCAGAATTCAGTAAGTTTACAGCTGAGGCC
TTATCTATAGACCATTGATTTTGCTCAAGGAAAAAGTTACACAACTAGCAATAGAGTCCTGACCAGGCATTACAAATT
CTACACTGATGTGTAAAAGAGGGACTAGGCACAAAGAATACATGCTTAGCACAACTATCTTTATTATAGGAAAAGCAA
TTTAAACATATTTTACTGAGTAGTGCCAGAAAATTACCGAAAAAGAAAGTTAATGCTTTTTCTTCTCAAAACCCCTTC
TATAATGTGTAGGCATTGTTCATATTAGAGAGACTCCTGGGAAATGCTTGGTCAACTAAAATTGTTAAAGAGCTAAAATT
GAACATTGACTCAGAAGCAATGTGAAATACATCTTCCCATTTCCAGGATGGAGTGCAGTGGTGGAGATCTCAGCTCACTG
CAACTTCTGCCTCCTGGGTTCAGCAATTCTCCTGCCTCAGCCTCCTGAGTAGCTAGGATTACAGGTGCCTGCCACCAT
GCCAGCTAATTTTTTAATATTTTTTAGTAGAGACGGGGTTTCGCCATGTTGGCCAGGCTGGTCTTGAACCTCCTGACCTC
AGGTAATCGACCTGCCTTGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCGCGCCCGGCTCACTGACGCTCTT
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ACGAGCCTCATTTCACGTCCTAATAGCAACATTTGAATGGTGGCCAGTGTAATGGAGAGTGCAGATCTAGAAGAACAAA
CACAACCTGGTAACAGAGTTACCTGGGGGAAGGTTGAGTTTGGGGATGGAGGGCTACAGAACTTTAGAGTTCTGCAGAA
CTTTTAACATTTTTTACAATGAGAATACATCATATATTATCTAGCTAATTTAAACAAATACATTGTTAAATGAAAAGC
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GTGCAATCTCGGCTCACTGCAACCTCCGCCTCCTAGGTCCAAGAGATTCTCATGCCTCAGCCTCCCAAATAGCTGGGAT
TACAGCATGTGCCACCATGCCAGCTAATTTTTGTATTTTAATAGAAACGGTTTTTCATTGTGTTGGCCAGTTTGGTCTC
AAATTCCTGACCTCAGGTGATTGGCCCAACTTGGCCTCCCAAAGTGCTGGGATTACAAGTGTGAGCCACTGCACTGGCC
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GCCTGTAACTATCTAAAAGCATAAAAAATAGTAAAACAAACGGCAAAATCATTTCTCTCTCTCTTTCTACCTGTGTC
TAGTGTCTATTGAACACTTACTATATTCCAGCATTTAAAGTACAAGAATCATGAGGCAACTGCTGTTTAGACTGAAGT
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AGTTGGCTTATGCAATGTTTATATAATTACAGGATATAAATGGTGGTTATTAGCCTAACTAGAATATATGCCTTTTATAA
TATTGGTGAATATTTCATATGTATTTTACTACAGCTTAAAATAATAATTTTCAGAAAGAAATTCAAATTGATTTTCATAA
TTTAGCAACCATTTGAGCTCTTCCAGGTCAGAATAACTTCTTGGGAAGCCCCAGAAAGAAAGCATATACCTGCTTTTA
CAGTTATCCATAGATTGACTATAAGGCTAGGTATTGAGTTGGCGGATGCATACTTTCTTAATTCTTTAGATAATAGGTC
AATAGTGTGGTAGTGAATCTTGATGGTAAGTGTCTTTCGCATGTTTCATACTGCACAGTAGACTAGACTGAAAGTCCCA
GGAGACTGTGAGTGTAGCAAAGAATGAATCTTGTATCCCCAGATACACAAAGACCATGTCTTCACTGGACCCACAGAAC
CTAGCACATGGTGTGTCTTATTAAATGCTTGTAGAATAATTAAATAAATTACATTTCAGAAAACAGTCTATAAGTCTTAA
AGTTTACCTTTCAATTATGTAATTATTACTTTTTAATTTATAATTTATTAACAGACATAGATATTCACAACACCTGCAAA
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ATCTTTTAGAGTGTATATGCGCTTTTGAAAAAAATGGTTTATGATTTTGTATGTAACTGTATATAAGTTATTTTA
GAAATTGCATGATTTATTATAATATCAGCAGTGAATTTAAACATTACACAAATGCCTGAATATAAAGAAATAGGATCAC
CTGTGTCCAAGTTAATTGCCTTCTTTTGCTGTATTTCATTAGCCATCAATACCTACATAATCTAAGGCCAAGATACATAT
GGAGCAATCTATGGACAGGCAATAGCATCCCATGGGAGCTTTGGAAATTCAGGAGCTCGAGCCTCACCCAGATGTACT

Fig. 9.177

Fig. 9.178

GATTCCTTAAACCAATTTCCCTGCCAGCTTCCCCAACAGCTGAGGAAAGAAAGGACCAGAAATGAGGGGAGGAAAATCA
TAGTTTGTCTCTGTTTTCTGCATGATCCCCACTTGGACAGAGACACATACAGTGTGTGTGTAATGGTGGTGTGTCTTCAT
AAAGCCCTTCCCTCACTCCTGACTTTGATATACAGTATTCATAATCTTAAACATGCCTTTCCAAAGGAAAGAAGAAAGGG
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AGACCATCATGTGAGCTTATTTGGAGTATTTGGAGTTTCTGCCTGAGTAGAGGCCATTCCAAACTCTGATCAGTGAGTC
CAAGTACTAGGGTCTATGTTACACACTTAGATAATTTTCAGGTAGTAGTAGGTTCTATACATTACATCTACATTTTGATA
AGATGAGTATAAATTATGATAAAGAAAACATAACAAAAAAGAATGAGTGTGTTAGTTGAGAACTGTATGACTAAATAA
TTTTTTACAGTTTTTGCCATAACTAACTAAACAGTATGTACGTTAGCAATGGGAGGGGAAAAAAGCAGAAACAAGGCT
TTTAGGAGGAAAGCAGCAAAAAGTAAAAGGAATTAAACCTAAACCAGTGACAAGACCGTGCTCAGGAATGCATTGATG
TTATAGATGAAAGTCAATGACAAAGGAAATAAAAGCCATGTACACCCAAAGACATGACCAACCTCAAAAATTTGGTTT
TAAGTAGCTAGATCTCTGACTGGGAAAGTGTTATTGTCCCCAGGATTTTAACACCTAGATGGCATTCTCTGAGTTTGGTA
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CTTTGAGGAAGATTTTCTTTTTGGAAAAAATTGTGAGCAACAAACAGTGAATAAAAGAAAGGCATGAATAAACACTGGG
AGATGCACTGAAGGAGAAGGTTGGCTCAAGGCCAGGCATCGGGTGCCAGAGCAGGAGAGCTGGCATGAGTTCCTACGGA
GCCAGGACCAGAATTTCCGTGTCCAGTGTCCCTTTTGGTTGCGCTGCTTTATCACTGCAGTTCCTAAATTGCACCCATT
GCTGAATGAGTGGCCCTCAACTCAGCTGTTTGATTTTCAGTCTGAGTGGAGCAACCTAGAGTTCAGCTGGAGGTTATCTG
AACTATTTAGACATGTCCGGATGTAAGGAAGGCAGGAAATCCCATGAAATAGGCTTTCTTCTGACATTTCTGGCACTT
CCTTTTCTGGTGCTGTTTGGAAAACTTCATGAGCACAATTTCTATCAATATTTTCATTTTAGTGTATCAGGTCACAGCC
CAGTAAGAAAAGCATAAAGAGAGCAATGGAAACCAGAAATAAAAATCATGGTATCGATTTTCAGTAACTCAGAAGGAAG
CAGTTTCCCTATGGACTAAACTATCAGGCAAGGGAGTTGGAAAGAGTATCTCAAGTTTCCATATTGGTTGAATGAG
CCCCAAGACGTTTCATAATATCCACAGGCATCTTTTGACAAAATCTCATTGTGCTTTCACAGTTTGTGCTAAGCACTA
TGGAGAATTTTGAAAATATGTTTCAAATTGTTCTGACATAAATAAATATATTTGTATCTAATTTGTTGCTAGTACAG
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AAATCAATAATATTGAAGACATCTGTTTTAGCTAAGATAATCTTGCTGTTCTACAAAATAGTCTCCAAATCTCATTGG
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GAAAGACCATGAAAGAGGATGGTATGTGTGAACCTTTATGAGACAGGCCGGGAAGTGCACATGTCACTTATACTCGTTT
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GAATTCCTAGAATAACATCTGTAATAACATATGCCATTGTTGAATTCCTAGAATAACATCTGTAATAACAAGTCCTTGA
ACCTCTCGTCAAACCTCCTTGATGAAGATTATACAGAAGCACTTCTGTTGTTTGATAGGGCTGTTGATTGGAAAGACCT
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AGCACTTTGGGAGGCTGAGGAGGGCGGAGTCCCTGAGCTCAGGAGTTTGAGACCAGCCTGGGCAACATGGCGAAACACC
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ATTTAGTAATTGTGTGACTTTGGTCAAGTTACTTAACGTGTGCTTCAGTTTCCCGTTAGTCAGTTTTCTGGGAATAGT
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GTCATTATAAGCTAGTATTTTTTTTTTATTATGTACTCTGTGCCATTATTCCAGCCTTGGTCTGCATGAACCTTTAGGAT
CATGTTTAAATTTCTAGATGATTACTCTGTCAAGTCAATTTGTGGATGAATTAGAGTGAAGAGACAAGAGTTGAAAGAAAC
GAGTGTATGTCAAGTTTTATTACTGTAGGAAAGGAGAAGGCAGAAATGAGAGATACTACTGAGGCAGAACTGTGGGGTTT
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CTTAACTTCCAGTACTAGTTTATTTTCAAGGAAATTGAAATCAGGAAGTTCTTCACATTCTTCAGGCTTTTCTCGTCATT
GAATTTTAAGCACAGTTTGGGGGTGTAAGGCCTAAATAAGTTTGAAGAACAAGGTAGAAATGCTGGTTTTTCAGTCTT
TGGGAACTTAAAGTTGCTGTTTAGTCATATTTAAGTCATAAAACCTGTTATCTTCACATATTCTTTTAAAGTAATT
TAGCATTATAATCACTTAAGTTATAATTTTTTTCATATTTTTTATAATACACATCTATATATCCACGACTCAGATTTTTT
TTATTTTTTATTTTATTATTATTATACTTTAAGTTTTAGGGTACATGTGCACAATGTGCAGGTTAGTTACATATGTATA
CATGTGCCATGCTGGTGTGCTGCACCCATTAACTCGTCATTTAGCATTAGGTATATCTCTTAAAGCTATCCCTACCCGC
TCCCTGACCCAGACTCAGATTTTTTAAATGCCAAAATTTTGTCAATTGTTTGCATCAGTTCTTTTATTTTTTAAAGAA

Fig. 9.180

TAAAATGTTACAGATAATGTCTAAAGCTGTCACCCATCCTATTCTTCTTCCTTCGCAGAGGATCCTGAAATCAGTGTGT
TCTTCCTTGAATGTCATTGTCTTTTAACTTCATATGTATGTGTCCTGTTTGTGTGTTTACCATTTTATATATAGAGATG
GTACATGTATATATATCTCCATAATATGCCATACTATAAAATGAGAAAGAGCAAGCTATATATATATATATATATATAT
ATATATATATATATATCCTTAAGCAGTTTGCTCTTTTTTTAGTACATAGTATAGGAATAGAGTTGTGAATATATATATAA
AATATGTGTGTGTATATATACATATATATGCATATATGTACCTATATACATATATTATATATACCTATATGAACCTATA
TACATATATACACATATATGAACCTATATACACATATACACATAYGTACCTATATACACATATACACATATGTGTACCC
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CCCATATACACATATACACATATGTGTACCCATATACACATATACACATATGTGTACCCATATACACATATACACATAT
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CATGTGTACCCATATACACATATACACATGTGTACCCATATACACATATACACATGTGTACCCATATACACATATATGC
ATATGTGTACCCATATACGCATATGTGTACCCATATACGCATATGTGTACCCATATACGCATATGTGTACCCATATACA
CATATACGCATATGTGTACCCATATACACATATACGCATATGTGTACCCATATACACATATACGCATATGTGTACCCAT
ATACATATATATACCTGTGTACCTATATATACACACACACATATATATATCTATATACCTACATATATATACACACA
TATATATATACCTGGATCATTTTTTTAAATGCTCAACAGTACACACATGTAAACAGCATTTTCAGTCAATGGTGGACTGCA
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CCTACTTATATACAAAAGTTAACTATAAAACAGCCTCAGGTAGGTCCCTCAGGAGGTATCTAGAAGAGGGCATTTGTTCT
CATAGGAGATGACAGCTCCATGCATGTTATTGCCCCAGAAGAGCTTCCAGTGGGACAAGATATGGAGGAGGAAGATAAT
GATACTGATGATCCTGTCCTTGTGTAAGCCTAGGCTAATGTGTGTTTGTGTCTTAGTTCCTAACAAAAATATTTAGAAA
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CCTGTAAGCTGCATTCGTGGTCACTGCCCATACAGGTATATCATTTTTTTATCTTTTATACTGTATTTTTTAAGTGTACC
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AGGTTTGTAGGCTAGGAGCAATAGGCTATACCATCTAGGTTTGTGTAAGTACACTCTAACATGTCTGCACAATGATGAA
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GGTGGAGGCGCTGCAAGTTGAGGGTGTCTGTACCATTTGAAAGGCATAGTTGGCTTTCCACGAAATCAAAGACCTGAG
GAAGATTTTCTTTTAAAGAACATGCCGATTGGCTTGTCTATGTTTAAAGGAATTACAGGGATTACATTATGACAATTTGGC
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TAAGAAAGTAAGGTAGATGCTACATGGGTAAAAACAGGAAACAGAAATTTAATAATTTCTGTGACTCATAAATAGGAT
TCAGGGCCTTCAGAATGAAGGTATTGGTGGTATTCACGTTAGGCCACCATCTGAAAGGCATAATTTTAGGTAACAGATA
GGGAAATGTATAGATCACTGTAAAGATTCTAATTTAAATTTCTCTTTTACTAGACTTCAATTTTATACCATCTTAACAC
ATCAGTCTCTTCTACTGTAAATAAGCAAAACAGAAAAATCATTTTATGTGCAGTTTTTAAGGACATAAATACTCTCCAAG
TATTTTCAGATGAGGCACTTATTTCTCAAAGGAGATCTTGAAAAGTTGATCTGAGAAGAAATTACATGATTTTCATTTTGG
AATAAAAACACTTAGGTTTATAATAATATAGCTTTTATAACCTTAGATTAAATTATAACATAGACGTCAATGATTTAGCTC

Fig. 9.181

Fig. 9.182

TCTGGACATTTCCACGTGGAAGATCCACTCTTAATTTATATTTAACACGCACAAAACCAAGCTTACTGTGCCCCCCTTT
TTGATTTTTCTACTTCTGTTTATCTGGCCATTATCTTCCCAAAAAACCAGAAAAAATCATATTTGACTTCTGCCTTGCC
CTCCTTCTGCCTAGGCCTATGGATTATTTACCTATTATCTCTTGCTGTTCTGTTCCCTCCCAATACCACCAGGTATTT
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TGTCTACTTATACTGTCTCATTAATTACCCTACAGTTCTGCTTTGATTGTCCACTATCTATAGCAGAGGTGAACAAAA
GGACCAGAAAAATAAATATTTTAGGCTTTGCGGACCCTATGGTCTCTGGGGCAACTATTCAACCCTACCACGGTAGTGCA
AAAGAAGCCCAGATATACAAATGGACATTAGCATGTTCCAATAAAACTGTATTTGCAAAAAACAAATAGGCTGAATTTTCG
CCCACTGGCCGTAATTTGCCGATCCCTGCCCTGTAGGATAAAGTTTACACCTTTAACATGGCATCTAAGCATCTGTCTAT
CAGTTCTGCACCTATCTCTGCAACCCTATGTGTTATTGTGTCTTATGGATACTTACATCCTTCGTTTTGAAGCCGAAT
CTAGAGCAGGGTTGCTCAACCTCAGCAGAACTGATGACTGGGCCAGAATGATGCTTTGTGGTGGGGGCTCTCCTGTGCA
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GGTTCAGGAATCTATTGCATAGTCCAACCTCTGCTACTCACTGTGGGATCCTAGGGTTATTTAAATTCCACAAATCACAA
TTAAGTTCATCTGTAAGTGGAACCTATTACTAATTCATACATGTTTCATAAAATTATTTAGAAGTTCAAATGATATATTT
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TGAAGCATGGGAGATTTTCAGCAGGGAGAGAACTGCAGTGAAGTGTGGAAGGAGGGGAAGTGACAGGAGGTGTTTTCTCA
CCAAGGGAATAGTGTGAACAAATATGGGGATTACATGGGGTTTAGGTGAGTTTGAGTGCCTCTAGGGAAGGTAGGTGGA
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GCACTGAACACTTTCAATAGTTGGACTTTGAGACTCAGTTCAATTAGACCAATCTATACCATCTTTCAGCTAGCTACTG
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GCATACAACACAAGTCGATCAGTAGAGTAGTGATGATACAGGACTGACTAGAGTGGAATATAACAAATTTCCCAATGC
TTAAACCAACAAGCATTTTATTATTGCATACAGTTTCTGAGAAATCAGGGAGCAGCTTAGCTGGCTGGTTCTGACTAAAG
TTCTTAGAAGAAATTGCACTCAAGTTGTGAGTGAAGGCTATAGTCATCTGAAGCTGTATCTGGGACTGGAAGATCAGTT
TTCTAAGATGGTGCTGTCCGCAGGCAGCCTTGCTTTACACTGGCTGTTGGCAGAAGACTTCAGTTCCTCACTGTTTGTG
CTTCTCCAAAGGGCTGGGTGTTCTTTTTTTTTTGTGTTGTTTAAATACTTTAAGTTCTAGGGAACATGTGCACAGCGTGC
AGGTTTGTACATATGTATACATGTGCCATGTTGGTGTGCTGCACCCATTATTAACCTCGTCATTTACATTAGGTATATC

Fig. 9.183

TCCTAATGCTATCCCTCTCCCCTCCCCCACCCACACAATAGGCCCCGGTGTGTGATGTCCCCCTTCCTGTGTCCAAGTG
TTCTCATTTGTTCAATTCACCTACAAGTGAGAACATGTGATATTTGGTTTTTTTGTCTTGCGATAGTTTGCTGAGAAT
GATGGTTTCCAGCTTCATCCATGTCCCTACAAAGCACGTGAACCTCATCCTTTTTTATGGCTGCATAGTATTTTCATGGTG
TATATGTGCCACATTTTCTTAATCCAGTCTATCATTGTTGGACATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATA
GTGCTGCAATAAACATACATGTGCATGTGTCTTTATAGCAGCATGATTTATAATCCTTTGGGTATGTACCCAGTAATGG
GATGGCTGGGTGTTCTACCTCAGGCACAGGAGTCAGTGATTCAAGAGAGTGCAAAGTAGAAGCCACAACGACTTTTAT
GACCCAGACTTGGCAGCAACACACTATTACTCTGCCATATTCTTCTGCTCACATGTGCCAACCCCTGGTACAGTGTGAGA
GAGGACTACTCAAAGTTGTGAGTATCTGGAGGTGAAGGCCCTCCTTATAAAAGGACTAAGTCAAGTGGATTCAAGAGATG
ATGGGAAAAGAAGTGGTGGCAATTGTTAACTCTTGGAGTTTGTGTGTAAGTCAAGTTGGAGGGGATAGGAAGAAAAATG
AGATATGGGCAAAGGGTGTGTTGTCAATTGTATTATGTTTGTGTGGTTTTTGAGAAAGAAGCAGCAGAAAGGGGACATTT
GATCATGAGAACTGGGGACAATTGCAGGAGCAATATTCTTGAGGAGGCTTTGGGGGAATAGGGGTAAGGAGAAGTTCA
GATAGTGCACAAGCCAAGGAGTTGCCCTTAATGGGAGTGCTGATTGTTTCAGTCTCACAATAAATGAGAAAGAAGAGTA
GGACAGTCTGAGAGTATAGGAGGTGAGTATCTTGATAGATCTGGTTTCAGAGTGTATGAAAAGTGTCTCTTAATTCCTT
CTATTTTGTGATAAAATAATAAGCTCATCAATGTAAAGACAGAGAACAGATGTTGGAGGTTTGAGGAGAGAGAAGGTGA
GAAATGATCCATCATCTCTGGGAGGAGAGAGTGCTTTGTCTAGAGAAATGTGAAATTCCCAGACGGCACTAAGGAGCCA
CCTCAGGCCAGTAGTTACAAATTCAAAGTCATCATGGTTGTGTTTTTCTCAACTATCATTGACTGCTTAGGCCACAAC
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AAGGTCAATGCACTGGAGGACTCATTGGGTTTGAAGATTTTTTGGAGTCAGAATACTGGAGGGGATGAGCTAGAAAGAG
AGAAGGTAGGGAGAGACGCTTAACAGTGTTTATGGGGCACCGTTATCTGTCTATGGCAAGGCCTTGCTTATGGTAAGGAA
AGGAGGTTTCAGGTAAGATTATCTGAAGGGAGATCCAGAACTGGGAGACCGTGGGAGGAAAAGATAATTATGAGTTAT
GTAAGAGTTTGCGCTGGAGAGAGACAGGGCTAAAATGTTTAAAGACTGAGAGGAGAGACCTTTAGGTCAGTAGAAAAC
GGATTAGGAGGGGAGCAGATTAATGATGTGAGTTACAAAGACATTTTAGGGCAAGGAAGAGATCATGGTCTGGAAGCAG
CAGAGAAGAATGAGAAAGACTCCTACTTCACTAGGGGCAGAGGTACAAACATGTGGAAGGAAAATAGCCACTGCATCA
GAGGGCTGAAGTAAATCCGTATTCTCGGTGGGCAGCCAGTAGTGTGAGAGCAAGATGGTGCAGGAAAGTCAGAGAACAT
GCTGAAGATACAGGAGCATAAGGAGATTTTGCTGATGACAGACTGAACGCTAGGAGTCACGCTGGCGAGTTTGCAGGAA
TTGGGGAAGGCTGGGGATTTCAAAGTAGGGGATTACAGAGCCATCTGGTGATGAGCAATGACCTGGCAGTCTTGGACT
TTGTGTGGTGACGGTTGTGTATTCAAGACAAGGACCTTGGTTTTAAGACCATGTAGTCAGACAAGCACAACTCCTCTG
CTGGGATGGCCGGTTTCTTTGGTCTATTTAATTAGTTCCAGAAACAGTAATTCATCCTCACTGCATTTATAGAAATTGA
GGGGAGAGGTGGTGGGTACCAGACTGGCCAGGTCTGCCAGTCATCTTCACTTCAACATTCCTGTCTGGTAAAGTCATA
TTATTCTCACATCTTGGCCGTGTGTGTATGCTCCAAGAATGAGACAGATGACAGAAGATACAGAAAAACAATTGGGAAA
TTGCTAGTTTGCAGATGTATAGCCCTGGCACAGATTTTCTAATATGAAATTGCCATTTTTTCCATGATTTGGACACAAA
ATCCTGTTCTTGAATATGATGTTTCAAGCTAGATATTGAAGAACTGAAAAATATTTTGAAAAAATATATATATTGCATT
TGTCACAACAGTTTATACACATAATTATCACAAATGATATTGCTTTTCTTAAATAAATAGGTTTTATGTGTTGGCATGAA
ATACTTTTACAAGATGATGCCAGGTTTTAACATTTATACAGGAATATTTTCATTATCATAATTGTTAATAAAACAAATA
TGAAATGTATAGCTTTCTCTTTGGAGTTTGAATTGTTGGTTCTTACTGCCTGCTTGCCATATTTATATTAGTTGTACTGG
CTGGCAGTTATTAGGAAATACGATCTTCTGTGCTTAGGAGTGCGCCCTGCAAGCAACATGTGCAAACATTTTTTTCTT
TCTTTCTTTTTTTTTTGGCCCTAAGGAAAAGGGTAAGCTCTGGCACCTTTTAACCTGTAAGGATTATGTGGCTCTTGCTGT
GATTGATTGGTTGGCTTCTCCTCAAGAAATGCTCCTGGTGCGTATTTTATGTTCTCACAGAGCCAGGGCTTTGCTGATC
AGCACGTCACCCCTGTCAAGACGTGGGCTTGCTTTTGTGCAATTTGCTCTTGACGCTGGGAAGCATATTTCAACACACT
GTCCAGAATTGGCTAGAACATGCCTGTCACTCCAGCTGACTCTGATGACGCTCTTGCCAACGTAGATTTACATCAACA
CAGTTCTTTACTGGAAAAAGCTCATTCAAGAAATACAGGGTGGCCCATTTAAAAGAGGCATAGCGTCAATTTGAAGAGA
AATATATTTTAAAGGGAGAACTCTAAGAAATACACAGTCAATTGAAGGCTCCATGGAGCATAGAATTATAACTAGCATC
ACCCGAGAATTTTTCCCCTGGAGATTGTCTAAGCTCTAACTTCTAGAGACTGGCTTATCCTGAAAAGAGAACCTTTCT
GTTCTAGACTTTTTCTTTTCAAGCATGCTAGATATGTAGGCATTTTTAGCCTCTTCTTTCCCTAACCATATAATCTGAACA
CCATATTCATTGAAGAAAGCATGAGATGTGAAAGCTGATAAGATGCAGGATTATGTGACACCATTGCTTTTTTACCTGTG
CAAAATTCAGTTTGGTAATCAATGGAAATACAGATTTTACCCTTCACATAAAATGTTTGGCATATGTCCATCTTGTTT
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AGGCATTCAAGAAATAAGCTTGTTACCATAGCTATCCAGGAAAAAATAACAACACTGTCTTTGCTGTACAAAAGAAAA
TTCAATACTACTGTACATGATATGTGATTACTATTATTATGACCCAATGCTACTCACCCCTTCATTTTCATCATTCTAG
TCTATTTTCAGGTCATTACAGTAATGGATACAACTGAATGGGTTTGTAAATTTTACCCCTTCTCTCATGACAACCAGCAGG
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GATTTTATGATTCTAATAGACCTCATGATGAGTTCTATATTTGTGCTTGTTTAGATACATCAAAGTCAGGCTTTATACT
AGGGCCACCTGGAAAGTTAAGGTCATCAGCTTTTTGTTATAGTCATTGGATTACTGTGATGCTCAGCAATAACAAAGTT
TGACCTTGTAAGTGATCAAAAGTTGTGGGTTACAATATAGGAATATTAAAGTATATTCTGCCATGATGTGTCTTCTGT
TGTAGTTTGTCTAGAATTAATATTACAGTCACAATTAACCTCAAAGTGTTCTATACTGCTTGCCATACAGTACAGTGGGA
AAGAAGAGATGTCAAAATCAGAGAAATAAGATTCTCCATTATGAATTAACATACAAAATCTACACTTAATGCATCAGA
ACTAAGATGCAATGTTTTAAAGAGATTGAGTTGACAGGTCAATCAGATGGTAGGACTGGTAAGAATTTGAGCAGGCATA
AACAATTGGCAGTATGTATTCAAACACTGTTTATGAGAGTGAGGAACTATTGGTTAAACAAGCTGACATTAAGCATTTAG
ATTATATAAATCACCTTATTAACCTGCATTCTATCTATGGATCTATTATCGATCTGTATCTATCTATCGATCTGTCTGTC
ATCTTTTGTGTTTGTCTGGTTCCATGCTTCCAAGACATTTTCTGATCTTCATTGTTACATAGTACTCTCTACTGAGTCC

Fig. 9.184

TAAAGAACACAGTTCTGTGTCAGGCTGATTGAAGTGTAATGATTTGAGTGATGATTGTAGTGCTCAATGAAAGGAAATA
AAGAATATTTGGACTCTGTTTCAGCAGTCATATGGGCTTGTAACAATATGAGTCTGCTTTTGAGAGAAGAACTTGCACAA
ATGTCTCTTTAAGAGCCTGCTTTTAATTCTTTTGGGATATATATCTAAGTGTTCTTAAGATAGTTCTATTTTAAATAT
ATTTTTCATACTGTTTTCCATAGTGGTTGCACCATTTTACAATCCTATCAACAGTGCACGAGGGTTCAATTTCTCCAT
ATTCTTGTCACACTTACGTGTGTGTGTGTGTTTACAGTAGCTATCCTAATGGGCATGAAATTATATCTTATTGAGGTT
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ATGTCTGTTCAAGTTATGTACTCATTTTTTTGGTGGAGTTATTTAATCTTTTGTTTCTAAAATGTTTTGAATAGATCAAG
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TACTTTACCATTTTTTTGGGGGCTCTTTTTTTTATATTTAACTTTTGTCTATCTGTAATGTGTTTTGATTATGAAGAAT
AAGGAAAAAATTTAGGGTTTGTTTAGCTTCCCAAACCCCACTTTATAGAATAATCACTGATCATGTACTAACTTATAAT
ACTTGATCATTTTAAGTAGGCTTTGTTATTGAGTTCCACATTTGGCACACCTACTGGTGTAATGATTGCTCTAGTATTA
GTGCAATAGTATCGTGTTAGATTTTGTGTTGCAGTGTGCTAAGTGGTAAACCTCAGCTATTAATCTACCTTGACTAAATA
TGTAATTGTTCCCTTTATTAAGCCCTATTTTCTTTCAAAATAAATATTAATCTATGTTTCCCACTCATTCTTTTTTGGGG
GAAGAGTAAATATAAAGAAATGTTTCCAGCAATGTGGGAGGCCAATGTGAAAGAATACTTCTATATTATAGAATGACTA
TTTTTGCAAATGCAAATGATAACAGAATTCCCAGATAAGGAATAGCAATGACTGTTTCAGGGCATTACCTAAGTGTGT
TTCTTGGGCTAAATTTAGAATGCCTGATTACTGCACACCTGGTATGAAAATAGAAAACCCTTGTCCTACATCACATAA
TGCTTACCACATTGAGGTACAGATAAAAAGAAGTTCTAGAATTAGGGTTGTTCAAAGCCTGGTTTACTATATTTTGGATA
TAAAATGGAGTAAATCTCATAATCCTTCTGAACATTGGTTTTCTCACTTGCTAAATAGGAATAAATAACATGCATGAAT
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CCTGTAATCCCAGCTACTCAGGAGGCTGAGGCAGGAAAATGCTTGAACCTGGGAGGTGGAGGTTGCAGTGAGCCGAGA
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TCTCAACAATATGTTTGCTCCAGAGAAGGTAGGGCCAGACTCCTAATCCCAATGTTGTTTAAAGCCCATTCAGATTAT
GAATTCATTTGAAGCATTTTTAGGACCCGAGAGACTACTAGAAGTTTACCGTTAACTTAGATGACATACAGTTAGAAAA
ATTTTCCATAGTCAGCTTTTTCTTCTCCTGTTATCTTCTGTATTTTACTAAGAGAATGTTTAGTTTAACTACTAGAAA
AACGTAATTTAGGTTGATTAAAAGAGAAGAACCTAGATATGTAAATAAAGATGAATCTTTTAGACATGTAGTCAGTAG
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CACCAAATGTCAATAGTACAATTATGGTCATATTGAGAGCACTTGTTATTAGACACACCCTTTAGATTTTTTTCTTTT
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CTAGATGATAAACTGAGGTTTCAGGATGGCACAGAAAAGTGTCCAAGGTCACACATTTAGGAAAATGTGGGACATAGAC
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TTATTGTTGTTCTCCTATGCTACTAGTTAGCTGAGTGATCACAGGCAAATTACTTAAATCCTCATAGGCTCAGTTTCC
TCTAATTTAAAGGAGAGTGTGGGGTAGTTGATCTCTAAAGTCCTCTTCAGTGCAAAGTGCTGTGCTTGCCCTCTTTA
CCTGAATAATTCCTAAGATTGCTGAAGCATCATTATCTCCTTGTTCTTAGCTACTGATTGTATATTTAATTCCACAATT
GGGTTTATCTCTATTTTTCATTTCATGCATCTTTGTATCATTAATTCCGAAGTCTCATATAAAATCTTTAATTCAAAAAAT
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TCCACACTCTTTTCCCTTTTAGAACAAAAATACCAGCAAGAAAAAAGGATGACTGTTCCAGGTCATCAAGAAAGTAG
ACTACTTGGTCATTATCAAAAATATTTTGAACATGTGGGCACTTTATAAAACAAAGAATAAACAATAGAAAATTATGTA
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TTATAGATGAATTTATTATTTCAGTTTCAGGACACCAGCAGCAATAAAGTATTTTCTACTTTTATAAATACTCATATGG
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TCTCTTCATATTGGACATTGGCTGTTTTTAATTGTTTTAAGATAAGCATACTTGAAGGTCAATCCTGGATTGGTGCATT
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GTTAGCAGTGTGCTTGTAGCGTATTACACTGCTGTATCGCTGTATCACTGCACCATCTCAAATATTGAGTTTTTTTATA
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GTTGCTGTTGTTGTTGTTGTTTTGACACAGAGTCTCATTCTGTCAACCAGGGTGAAGTGCAGTGGCACAATCTTGGCTC
ATTGCAACCTCTGCCTTTTTGGGTTCAAGCGATCCTCTCATCTCAGCCTCCCGAATAGCTGGGACTACAGGTGTGCACCA
CCACACCCAGCTAATTTTTGTATTTTTTAGTAAAAATGGGGTTTACCATATTGCGTAGGTGAGTCTCGAACTCCTCACT
TCAAGAGATCTACCTGTCTTGGCCTCCCAAAGTGCTGGGATTACAGGTGTGAGCCACTGCACCCAGCCCTTTCTCCCTC
ACACTTTTGGAAAAAATATTTTTCTCAGTTTGTGCTTGCCATCTAATTTATTACATTGCTTTTTTAATATACAATA

Fig. 9.185

GACAAAAGAATCACTTTTCTTTTGGAAAATTTGATCTATGACTTTCATTTCATATAAAAAGCTTTCCTCACCCCTCAGGTCA
ATGTTACCTTTTATTTTTTTCTAGTTTTTTTAAACTTTGTTTTTAATCACTGTGAAAATAATTGAAAGATGATATTTA
GATAAAAATTTGTGTAAAGCTCTACTATTTTGATACAGATTGAAAAGGATAAAAAGAAATGAGTAAGAAGTATGCGTAG
TGTTTAGAATGAACATGAATGATAAACTAATAGGTATTATTTTACAAATTATAGAATATTAAAGATAAATATACTCTAG
AGTATAGAAATATATTTTATATTATTATGTTAAATATTTAAATATTTTCAGGCATGTAAAAATAGATGATAGTGTAGCCA
TTTATGCATCAAAGAGTTCAAACCATAACAACACTAAAGGTAAAACCTGGGGCCCCCTAGTGTACCCTTCCCCAATTCTATC
ACCTTTCCTTCCCACCCAGAGGTAACACTATTATGTGTTTACTTTTCTTGAGTATGTTCTTGAATAGGTGTGTATTCA
AAAGAGACAGTTCCTTTTTGCACTTTGGAAAACAAATAATTCTGCCTTCCCCCCCATGATTGCATTTTATATGTTTTAC
TAAGATATTTATTATAATAAATATTACACCACCTTTTCTCTGTCTTATTCATGGGGAAAGGAACCTTTTTTGTTCAAAAT
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TACAAATATGTCAGCTTATGTATGCTGTTGCATGTGTTGCTCAGCATTGCGTGTATTACTGAGCATTGCATGTATCACT
TAGCCTGTCTATAAAAAAATTGAGACATTTATTTAAATAATAATTGTCCCAATCTATTTCATGTTTATGCATGTGGATA
TTTGAAGCAAAATATGCCATAGAAACAGTATCTCTTGCTCACTGAATTTTAAAGTATTTTAAACATCAGTAGCAAGTGCA
TAATAAATCTTTAATGTATAATACGAACCTGATAGAAAAATAGGGCTCTGCATATTTTTTCTTGAAAAACAAATGTGTTT
TGAGAAATATAGCTTTGTCTAATGCTGCCTTTGTAAAACCGGGAAGTCAAAGACAAATAATATTTTGATTCTATATGT
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TTCAGATCTTTGCAAAATCTAAATGGGCTTGTTTTTAATTTTCATGTGGACATTTGAAATTTTATAGATAAGACTAACA
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TCACAATACCGTGGCCTATTAGATTGCTAAGAAATTATGTATGTCAACTCTTTCTTTATACAAATATTGAAAATGATAG
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CACTTCAGGAATGCGACTTTAAAAGCCTAGGCTTTATGTTATGTCAATCAAGTCATTGTACTCAATTAGACAATAGTG
CTATAATTAATTAATGTTATTATGTAAATGCAGAATTACCAAAGAGAATTTCAAAGAGTATAACAGTATTAAGTCGTT
TCCTTAGTAATTTCTCATTTGCTGTGAAAAACATCTGTAACCTAATACATTTATATTCATTTTTTCATATTATATTATGCA
CATATTATTTTTTAAATATTGTTAACAGGTAATTTTCTGATTACCAAAGTAATACTTTTGTATTAAATGTTTGAAGACT
GCCTCCAAAATTTGCAAAGAAGAAAATTTTAAAACCGTTATAATCCTACCTTTGAAATAATTATTCTTCAAATGTTAGA
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CAAGCAATTCTCGTGCCTCAGCCTTCCGAGTACCTAGGATTACAGGTGTGCACCACCATGCCCGGCTAGTTTTTGAATT
TTTAGTAGAGACAGTGTTCGCCATGTTGGTCAAGGCTGGTCTCAAACCTCTGACCTCAGGTGATCCACCCGCCTCAGCC
TCCCAAAGTGCTGGGATTATAAGAGTGTGCCACTGCACCCGGCCAATTGTAAGAATTATTTTCAAAGGAATTTATATCA
AGTTACAGTGCCCCAGAATATCTGTTATTTTAGCTGTATTGAATATCATAATTTTCTTAACATGTTTTGTCTTTAGAT
GGTATAAAATACTATCTCAATATTATTTTCCATTTGTTTGTGTTTACCCTTTATATTTCCATTTGCTTGAACCTCTCTGCTG
GAATCTTTCAATATCTACCTGATGAATCAATGTTTATCTGTTGGTTTGTGTTTTCTTTTTTTTTTAATTAATTAATTTATTA
TTATTATACTTTAAGTTTTAGGGTACATGTGCACAATGTTATACATGATTTTTATTTCTTTTTTGTACATAAATTTGAC
TCTTTATTTTAAAGCAACTATTTCTAGTCTTTTGTGTTCTCTGTTTTATTAAATTTTTATTGTAAAGAAGTTTTGTTTTG
GCCAGGCATGGGGTGGCTCACACCTGTAATCCTAGAACCTTTGGGAGGCCGAGGCAGATGGATCATTGAGACGAGGAGT
TCAAGATCAGCCTGACCAACATGGTGAAATCCCGTCTGTACTAAAAATATTTAAAAAATTATCTGGGTGTGGTGGTACAT
GCCTGTAATCTCAGTTACTCAGGAGGCTGAGGCAGGAGAATCACTTGAACCAGGGAGGTGGAGGTTTCAGTGAGCTGAG
ATCGGGCCACTGCACCTCCAGCCTGGGCCATGAAGCGAGATTCTGTCTCGAAAAAAAAGTTTTGTTTCTGTGATAGAT
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AAGATATGTAATCTCAGTGTATTATATAAAATGTACCATTCTCACTTTCCTTATTTTGTAAAGCTTGTTTAATTTAGAG
TTTGATATATTGAAGAGTCCTATTCCACTGACTTAAAAAATTGAGGCAGAAGGATCCCCCTCAAGTGTCACATCTTAGA
ATTTGTTTGATGGTACTTCAAGGCAGTTGGTGAATAATTTAGAACTCAAACCTTTGGGCTGCAGATTGCCTGAATACAAT
TAATAGAAAACAAAATATTTCTCAAATTACATTTCTTTGACATTAGTAATCATTCCTTTATATACATCTCAAGTCTAAA
CTCCCAATCTGTTTATATGCAGAGATTACAGCTTTAAGATTTATGTTTCATAACTGCAATATCACTCTATGATACATT
AATGGGATTCTGTACTCAACTATTTCCATTGGCATTCAAGTGAATAATTTTTATACAAAACCTTCTTCAGGAGACAGGCCC
AACTGAAGTGTATCACTTTAAAACAAATATCCTATGGGCAATAGATAAATCTGATATTTTTCTGAGTAGAAGAAACATA
AAACCTCAATATAGGATTAATAGGGTTCAAGGGGTTTTATAAGCACAGTGCTTGTGAAAGTATGTAATTCCTATTAAGG
CTTGCAATTCATGAGCACATCATGGTATATGCTCTCTCTGGGAATATGTAAAGCCAGTTTAAATTCATTACAGACATT
GCAGTCTAGACCTCCGCAAGCCTTTGCACGTGACTGGACTCACTCAGGTAAATCTCTGAACCAGTTGTAGAGAGCTTTT
TTCATCTCTCCCTCACACATTTCCAAACTCTTTTCTTCTGTGTTTTTTAACTGTGACTTTTCTTTACAAAAGGAAAAG
AGAATTTTTTTTTAAGTCCATGACATATCCAATGAAATCGAAATGATTTAATACATGGAGTTATCTTAATATCTTTTGCT
TCTTGACAGTTCTCTCATATCTATCTAGAAAATATCTGGCCAAAAAAACCCACTTTACTTTGTTTATGAGATATTAGAT
TATTTTGTACATTTTCACATTCCAGGGCTACAAGGAGACCACAGTAGACAAAATCAAACCGTCTTCTTTCTTTATAG
GCTTGTCTCAGCAATGCAGCACTGTAATGTCTCTTACTTGAAGGAACCTTCAATCTATTGTAAATTCCTAGGACAAATAG
AATAAGGAGATAGAAGAGTTGTGTGAGTTATAACTTATATGTAATTTTCTGTATATATTTGGAAAGTTCAACATCAGAT
ATTGAGTCTATTTCAGTCTGTGCTGTGAATAGAATACACTATTTTATCCTATTGGTTCTTTTCTTAAACTGTCAACCA

Fig. 9.186

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TGGTTTGAGCATAGTTTAAGAAAAAAAAAATGAATGCTTTAAATAAAACCAGTACCTGGGTTGGAGGTGGCAAATATAT
ACAAACCTGTGTATGTGTGTTTGGGTGTGCGTGTAACTCTGAGGATTATAGCCCTGTATCTGTTTCAGGGAGAATTTTTT
TTACATCAGACCATGAAATTGAAAGGTGATTCTGATAATCTATTTTCATTAAATAGAAGTTTATGATACCATAGACTCTG
GAATAAAAAATTCCTTAAATTCCTTATTTAGATTAAATGCAGATTTAATCCTTAAACCATTTGGTTCAAACTCAAGTTT
ACTCTTTTCAGTAATAACAAGAAGTGGTTGTCAATAATACTCTCATTAAAAATAATTATTTTCAGCATTAAAAAGTAATA
AAATTGGTATTTTCTAACTTATATGCTTAATACTCATCCACAAAGGTTAAATAATTAAGAAATTAAGACTGTGAAGAA
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ATGTTGACATTTCAATTTTTTGAATCTTGGTAAAATAGTCATTTTAATAACATACTTACCAGATTACTTAAAGTAACAT
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TTTTCTGAACTTTCATTGGCCTTGGATTTTGACCTTCCATATACTTGAAATTTCTCTCTTTTTTAATATCACCACCAATAA
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ATACCATCAATTCTCAATTATTCATGGGGTTATTATTCACAACCTTACTTGTGATATGTCCTTAAAGGTCATATTTACAT
AATGAATGATTCTGCCTATTTTCATTGGCATAAATAGGTTTCATTTAAACAGATTTTCATTATTTAATTAGCTCCAATTTTT
CCTTTACCATCTTTTTTCAGAAGGCAAGATTTTGCTTTTAGCTGTCTTGCTTAAATCTGAAAGCTCATACTGATCATTTCT
CATTTAATAAGGGAAGATGAAAAGGGCCGCTCAGGGGACTGGGATCCAAAAGGTCAAGAGTTTTCCATTCTTCTTCCA
TCAGATCTTCATAGTGGAGTTTTGTTAATTTGTGAATTTAGTTCAATGAGCCAAAAAAACCTTGCTGATCAAAACAAA
GCCCCTTTGGTAGGGAGGAAATTTCTGTCTCTTTGTTTTCTCTTCTCTCTGTGCGTGAGTGAGTGTGTGTGTGTGTG
TGTGTGTGTGTGTGTGTGTGAATGTCTCCTGACATTTGAAATTCACAGTGCAATTTAACAATAAAATAATAAATATAG
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GGAAGTCAATGATGTTTGCAGCAGTGTATCTATCTAGACAGATTTTGGAAGATGTCAGCAATTTTGTTTAGTCTGTC
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CAGGGTAAAAGAAGAGAAATGGAAAGGAAGTAGGAAGGGATGGTGTCTGTATCAGGGAAACAAGGATTTCTTAGGAATC
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CTACCTTCCAGGGAAGCTGAGGAGCTTGTTTTTTTTTAACTAGACATATTATTGGGTGATTAAACATTAGGCTTATATAAA
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CAGATGATATTACCTGTATACTATTATGTTAGGGGTATAAATACCATAAATATGCTAAATTCAGATTAATAAAAAGTTTT
TCACTTTCTGCACAATCATGACCTCCTTTTATTAAAAATAGACAAATAGTCTGGGGTTAGTCACACAAAGTCTATATGGC
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CCTGGAAGCCACAGAAGATGCCAAATGGACACATTTTGTGACTATTTATAGCTGATCCAAAAATGAAAGGGGAGAATGA
GGAAACTGATTGATAGTCCCCAATTCTGTTTTGATCTGAAATAGCTTGTTAAATGAATCTCATTTGTCAGAGCTAGTTTA
AATAATAAATGTCTCTACTCTCCACTATTCCGTGCATATGTGAGTCTGAAAGTAAAAGTTTGTAATTAAGTGACTGCA
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CTTTCTTCCCCATTCCAGCTTGACATTCCTGAAGTTTTTCCGGCTTGGAAACAAATTATATTACTAGGTCTGCAGTATCA
TTTAGATTGAGAAGTATTGTGATAGAGCAAGGTTCAACCCTAAGTATTGACTCTTAGAATCTGCTCATTCCATAGTCTGT
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ACTCACAATTAGGTTTTCTATTCTGGGGCTATTTTGTACATTTGTGTACAAAAGACTGCATTTTTAAGGTGTCTGACTTT
TGACTTGGTTTCAGAAGGCATAAGAAGTTGGTTTCATGGTTACAACCCCATGCCCAGAAAAGTTAAGAGTGGTATTTTT
AATTAATTTGTAACCTAGAATTAAATATAGCTAGAATTATATAGGAATACAACAAAAGATTAAGATGTATATACATCCCT
CTGAATAATATAAATATTGACAGGCTGATATTGGGTTTTCTAGACTTACTTTTCGTAGGTATTTTCTTAGTATGAAGATA
GTTAAATATACTAAGTGAAAAAAATACTCAAGTCACTGTATTAGTGGTTCCACTAGTAAACTACAGACCTCATCT
GTTTCTGGAATTTGCAATTTTAAATATTGATTTTTGGAATGTGGCGACTTACACTGCCTACTTTTAGTTTGCCAGGTGA
GCTCATTTTTGAAAATATCCTATACTCTGCATTTATCCTTTGATAAAATCATGGCTATTATAAATTTTTAAAAATGGTAA

Fig. 9.187

AGATCATGTCAGAATAATGAATCGTGCTTCTAATTTGATAAATGTAGCTTAATCACAAATATACGAAACTTTCTGTTGG
TCTGGTGCAAACCTTTCTCAAGGAGAAGATTTAGTGTTTCAGAACCATTGGAGTAGATGCTATATTAAACATTGAGGCTC
TCTCACCTTTAACTTTATAAAGCTAAATATAGTCAACAAATGAATTGGAAAGACATATTAGTAAAATCTACCGAGTCTA
GGCAGGCCCTATGTGTCAGGCCAAGATGCTTCCTGGAATTATTCTGTTTCAGAATGATTCCCTCTTTCTTTCTCATTTCTCTA
AAGATAGGATGTCAACAGTTTCTTCATAAATGTAAGGTTATAAACCATTGACATTTGTGAAACTATTGCAATAATCTTGG
GTTGTGTAGGGATAGGATGACATTTTCAGAGGAAATTCTCTAAACACTTTTCACATTAATAACATCTGGGAGCCATAATA
TTATAAGCATAAGAAGTGGATCTCATATAAAAACATAAAATATTTTTTTTCACATTTATATGAAATAACATATGTAATTG
TAACAACTGAGAGTCCCTGATCAAATTAGGAAATCTACAGCCATTTATTTTAAATCTTGCTCTACCAAAATTGGATTGA
GAGACAAATTAAACACAAATCATTAGTTACATTTTATAACTATAGAGGATGTGCAGAAAACAAATATTCAAACCCCAATT
TTATGTACTATATTGTGCCATATAGTCACAGGAAATCTGTTTTCTTCTAAATTATTAATAATTTAAATCATGAATAA
TTGTTTTATGTAAGACAGCTGGCCAGGCTATTAAGGGAGTACAGCTACACAGTAACACCAAGAGTGAGGTGTCCATCTGA
GCTACCACATGGAGAAAGAGCAGCTGGCTTTGCCCCACCATATTCAGTTGAAAGCTGCTTATTACTGAGCATTTGAATA
ACTTGAAAGGAGAGAAAAGGAAAGAACAACGTCAGGGTAGAATGTTTACTTGTTTGAGCCTCCAGGAGAAATGCCATA
CAAAGATTTGACTTGATTTTTTAAATCAAACCAAAGATCTTTCAGATTACATTGGAGGGCTGAATTGAAAGTTAACATTG
TATAAGTTGTATTATTTTTTATGTTCTTTTGTATTATACTATGTTCCCTTTTATATTATGTTTCATATTACCATGAACCTT
AGAGTCGGGATAGATCAGAGCTCAAGTCTTAATGTCATCTCTCCAAACTGTGTGACCTGGGAAAGACATTTAACGTGT
CTAAGCCTCAATTTCCCTCATCTATCAAATGGGGATAATATGTTTACTGACTACAAATTAAGTAAAATAAGACCTACAAA
ACTGGGTGCTACTGTATGATAACACAACAGTTACTACAGATGCTATTATTATTAGGTACGTGTGTGAAGAGAAAGTAA
AAAGAAGACAAGATGCAAAGATAAAGAAGGCCAAATGCAAAGATCCTATTTCAATGAAGCCAATAGTAGTCTTTCAAGT
TGCTGTTCCCTTCTTATTTCTCTTCCCTTCCAATAATAATAATTCCTATTGTCTACCAAGTATTGAGAGCCTGTCATAGGT
CAGACACATGGCATGCATTGAGAGAAAGAGAATTCAAATTATTCTCTTGAACCTTTTGAGCAACTCTATCAATGGGTCTT
ATTTGGTTCCATGTTACTGCTGAGTAAAAGAACATGCTCACAGAGGGTAAGAAACCTGCCACCAGCACCTTACCAGTA
GACATTAGAGTCAGGACTTTAAACAGGCTTTTGAGACTCCAGAAACCATGCTCTGCCTACTCAAATTTAGTAGCTGTTCT
GAGGATCTCATTCTCACTTATTTCCCTTTTTTATTTGTTTGTATTATGTTCTAACCCTGATTTGAAACAAATTTCTCAG
ACCAGAAATTTTTACTCCTGTGGAAAGAGTAATTGGATGTTCTCATTACAATATTTTTTTCTTCATGAATGTAATCAAA
CATCAGCTGGAAGCTTGCTTTTAATGCGTGGATAAACCAGCCCACACATTTTATTTTCCCTGAGGCGTATTGCTGGGAA
AATCTATTTCTGATTTTAAGAAATCAACATATGCTGAACCAACTGTCTATTTTCAATTAAATTGACAGGAGCTGAGACT
CAAAAATTATGTTGTATTTTCCACTTCTAACATTGTTGATAGAAAAGTTGAGGTAAACACTTTATTCTGCCTTAGATGC
CTGCTAAATTAGTCTCTCTAAGTAAAAATGTCAGGAGGCTATTGTTGGATTAAATTTAGGTGGAGTTGATACAGTTTTAG
GGAAGAACCGTTCTACAGTATGAGAACAACCTTTTAATATTATGTTAGTTATAATCTGGAGCATTATCATACCTTATTA
TATATCGTATGTTTTGACTTTCTCCTCTATTGGGCTGTGGGCTCAATTAAAGATAAGAATGTATCTCCCTTATCTTTGTA
TATCGGTGCTTAGTGCAATTGCCTGGCACATAGCAGGCGCTCAACTCTGTTGAATTAAATGATCAATGAGTTAAACTAGA
TGGCTATATAGTTGATCTTCTATATAGTTGATCTTCTATATTCAAATATAGTTTACATATATGTATGCAAGTGTATAT
GTGTGTTTATGAATAAGAAGAACTATATCTTCATACTGTTGAAAAGTTATATTGTTTCTCATAACAGATTTTATTCCTTT
AGCTAAAGTTGTATCAATATTAAATTTATAACTAAAACGATATTACGTTGGCACATGAGGGTTTTAGTCAGCTCCAGCT
GCTACAACAAAATACTATAATCTGGATAACTTACACAACAACAGATTAATTTTCTCACAATTCTGGAGGCTGAAAGTTC
TGAAATTAGGGTGCCAGCATGTTTAGGTTCTATTGGGGGCCCTCTTCCCTGGCTTGCAAGTCAGTGCCTTCTGACATGGTG
GAGAAAGAGCAAGCTCTCTGGTGTCTTTTCTTATAAGAGCACCAATCCCATCATTGCAGGCCCCACCCTCATGATCTCA
TCACCTCCCAAAGACCCATCTCCGAATACTATTACATAAAGGGTTAGGGCTTCCACAAGTGAATTTTGAAGGGGACAT
AATTCAGTGCATAGCAATGAGTATCTTGTAGGCTTATGACCATATAATTTGAAGCTATGATTTATGTAGAAGTTGGACA
AAAATGTTTCACATAATTAGTTATGTTACATGATCTAATGCCTGAAACTCATTTTTTAAATATGCCTGAAACTCATTTG
TTAAATTTATAGTGTTGGGAAAAGAAGCATTTAATATTTGTTAGCATTTGAACTTAAAAAAAATTAACACATTTGACTTT
AGAGAAATGTATATGTTGAAGAGTAGGTGTGGATGAAATGAAAGAAGAGAGCTAACCCTTCTCAGTGCCAGACACAGTG
GTGTGGGCATTAATAATTCAACCCTCATAGCAGTCTTAGGGTGGTCTTAACCATTTTCATGGGGAAATTGAGACTCAGA
AGTTTACTAATTTGGCCAAAGTCACATAATTGTTTAGAGACTGATTCAAATCAATGTCTTCTTTATTCCAATCAATCTT
TTAGGGACCTTATCTGTTATTTATATGAAAAAAAACAATACTTTAATGTTTATGGAGTTTGTTTAATTAAATAAAGC
CCTTAGGATTTCAAATTGTTTCTAAGTCCAAGAGATCCCAATGAGAAGGAACACAAAGTAATTTTTAATGATTTAAAG
TAAATGCTTTAGACTGAAATGGTAGCTTCTTTTTTAAATAACATTTCAAATTAGGTTTACTATTAACATTTACAAAGAAC
CAGGGCTATAAATCAACTTTATATGCAGTGCATTCCATTGATCTTAATAGTTGTATTTGCAAGCTGAGTAGAAGAGATT
TACAGTTCTCTCTCAAGGGATAGATCTCCTTTTTTGGGCAATGAATATACTCCTTCAGAAAGTCCCTCTTGCTCTTTTAC
GTACTCTTTTTTTGCTTGGTATTTCAAAGGCTTTTTTGTATATACTTTAAAAATATTCAACAAAAAGTCACTGAATATCTAC
AATGTGTTTATATCTTGGGTGTAATGCCTAGGTTAATAAGATGCATAGAAATATGGACCCAGTGCTACAGGCCCAGCAG
TGAGCAAAACTCAGGGAAACAAGTGACACTAAACAGTGTGGAGAATGCAAAGGCTAAAGGCTATATGAGTCCTGGCATG
AGCACGGAGGTATGGGCTGAGCAGCGCTGCCTGAGGAGGTGATGTACAAGTGGGTATTTATCCCCCGGGCAAACCAG
AAGGAAAGGTAAAGCCAAGGAACAAAGTAGACAATGGAGATGGTGGGGAATGGAGGGGAGAAGAGGAAGAATCTTGGAG
AGATAGTCAGTCTTAAGGTCCAATTAAACTATTTTCATGGCAGCCATTATTATCCTGAATTTTATGGATGAAGAAGTGAG
GTTCAAAAATATTACATAACTTGCCCTGTGTGACAAAATAAATGAAAAGCAGAGCCAGGTTCAATGAAAGTATGTTTGA
CTTAGGTTTTTAAATTTCTAGTCATAAATGGCAATTTAATGTTGTTATATATTTGTATTGGGCTTCAGCAAAAACAAAAA
TAAAACCTCAAGTACACAAAAGCTCTAGTAATAGAGCCATGTTTGTGCAGTTATTTCCAGCAATCCTTGGAACCTCCAA
AATTCTCCTCAGCCTGACATTATATGCCCTATGTGTCATTTTCATTATGTCACCACCTGGTGGTGCCTCACACCCACTT
TGGGAGAAAGGGGGTTAGAGAAGGGAGGACATCTGTGGACAAGCCAATAAAAGCATTTACTGGCATCGTGACTTCATAA

Fig. 9.188

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TTTACACTTTTATTTATGCATAATTAAGTATAAATAAACAACATGTATTTTCTAACTGCTACCTGTTCTGTTTCCA
GTCTTGCTTGAAAATCATCTTTCTCAAAAACTACCTATCATGTGCTATGCCCATTACCTGGGTGACAAAATGATCTGT
ACACCCCTACAAAATGCAACTTAGTCATGTAATAAAGCTGCTTATGTTCCCTCTAAACCTAAAATAAAGGTTGGAAAGG
AAAAATAAATAAGATAAAAAATTATCTTTCTCTATCAGAGTAATTTGACATCTTGAGGAAGTGATCCTGGGACTTCATA
TTCTTTAGGATTCAGGTGTCCAGATAATCCCAGAAGTAGCCAGCAATTTGGCCATTTGGGGTGTAGAAACCTTCATACC
TAGGTTATTGTAATACCTCAGGCTATTCTCTCCAGGTTTCCAGTGATTAATTAGCTGTGAACCTCAACCAACACCTGAC
TCTAGAAGAGTCTTCAAATAGGTCCTAGAAGGACCAAGTAAATCACCATCCCTCAATCCCTCAATTTTTCTTTTCC
AAGTCAAGAATAAAAGAAGTCTGGGAAACATTGCCAGGTGAGCTTCTTTTAAAGCTCATGATTTTCTGCTACCTGAGGG
AAGGAGAGAAGGAAAAAGAAAAGAAAAGAAAACCTCAATGAATGCTCCATAACCTGGATTTTAAATCTCTCTTTCCCTTT
TTGGGATAAAATGTTTAAATGTAATTAACCTACAAGGAGAAAAGTTAACCAGTGGCTTCTGCTTTTGCTGAAAGCACTTT
TTCAAACCCAGCTGTCATGTCAAATGCATTCAATATTAGTTTGGACAACCTCTTACGTGGGTCTCAGAATGCATCTCTA
AAGAAAGTGTTTAAATATTTTTTAAATGTGAAAACCCATATGGGTATTTTGGTGATAGGATTTCTTCTATGATTCAGGAA
TAAAGTATAATGCCCAAATAGGCCCTTGCCATTCCATATCAGGGACATTGCATCCATAATCCATTTTCCATATCCGTTTG
TATGTAAATGAAAAGTCTCACACATCACAACCTTCTGTTTTTCTCATTTGTAGGATCGCCTCATCTGTATTTATCCACATT
AGTAAAAATTATCCAAATCCCAGAAATATAACAATAGTCCGTATTTCTTGAGCATTTACTCTGTTCTAGGCTCTGTGTT
AAATGCTTTGCATGTAATGTCTCATTTAGTTTTTACCACATTCTATGAGGTATTTACTCTCCTTTTCTTATATGGGTAAG
AAAACATGTTGAGTTATTCAGGTAATTTGCCCTATATTCACATACTGGTCATGAGAAAACCTAGAGGACCCAGGAGATTGA
AGCCCATATTTTCATCATTTGACTGTCTTAGGAAACGCCTTGATATACTTGTTTTTTTAAAGGTATTTCTGTCACCCAAGAAC
ATTGAAGGTATATGCAGATTCTCTTTTCTGTTCTATTTCCACTAGACCTGAATATGAGGGATGAAAATTGCTTTGGTTT
TGATGTGACTTCAGTACCTTCTGTATTTGACACAGAGGTGAGTCACATCCTGATCAGTGTGAAAGCATTTAGTAAGAT
TATTAGTTTATAAAGAAGGCTGAACCATGACTATATAATAATGAAGCAATTGTAAAAAAATCAGAAAGCATTCACATTC
ATTCCTTGAATTTCCAAAAGCATTCACAACAGAAAAAGGGCAAACACGTTTATTTCCATGTATATATACTGAATTCCTA
CTATGCACCTTCAACAATAAAGGCTTTTGGTATATAGGAGTATACTGAGCCCAGTTAGATTCAAGTGAAATTCCTAGTA
ATGTCTTGGTATACAAGGAGCTTAAATGTTCAACATATAACATTGTTTTCTTATGATAGGTTTCACCTATAAGCCCTCTA
AACTGTTTGTAACCTCTATAAGAAATAATTAATAAACCCCAACAAGGTCTATTTAATGTCTTGAGACAGATCATTCTACT
CACTTTAGATTTCTTATAGATAATAACCAGCACCTAAACAATTTTTTAAAGACATAGGAAATCCAGAATAAATATAACT
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GACAATGAAGATTTGCAAAGGTTCTAGAATACTTTAAATTAAGTTAGAATAAACTTTTAGTTGCACTGTGCTTTGACTT
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GATTGAATGAAGATTAATTATAGGGTATGGGAATATTCAAACCTTTTTAATTGTTCTGAGTAGTGTCTTCTGCTGTTTT
GTTATCCAAAAGGGAGTAAGTATTTGGGGAACAAAGATTGTGACACATCTGGTAATATTCAAGATGCACACCCCTCAC
TAGACTGTCAAAAGGCTGGGCTTGTCACAGATGTGAGGCTGTGATGTATTGTCACTCTTGCTGCACCCATGGATGCGC
CTCTCCTGATATGCGACTCCCATTTGGGAATGTGAAGACACAGAGGAGATTTTTAATCAAGAGCATAGGTAGGAGTCAAT
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CTTTTATGCAGGACTCTGACCTAATTATTTCAAAAATTATTGCTACTGATTATAATCTCATTTGTTGCTTTCTTTTATCT
ATTCATTGTTTAAACAGCATTTTTTAGGGGGCAGGGTAGGGAGAAGCAGTGTAGCTTGGTAATAAAGGGCATATGATCTG
GACCCAGCCTACCTTGAAAATGATTAAGCCCCAATTTTCTTATATGTAAATGGAGACAATACTAGCATCTACCTCAC
AATATTGTTGATTGAATGAAATGAGATAATATAAGTAAATTACTTCCAGTAGTCCCTAGCACATAAGCACTCATTAAT
GTTAGCTTTTAAATTTGTATTTCCAGATTAACATGCCTAAAACTAGGGCTACGTACTGAGAATTCATAGAACCAATTTT
ACTTTGTGTCTGTGTTCTTATGCCTGCATCAGGCATTACGAAGAATACAGAGAGACAACTTTCTATTCTCAGGGTTG
ACAATFAGAAGAGAAATAATAACACAGGAAATAAGTAAAAGTATAATTCAAACCTCAAGTTTAAATGATGTATAAAATG
AAGAGGTCAGTTTGGGCTATAACAATCAGATAAGATTTACTAGGTGCTAAACTCGATCTAAACTGAGAAAAATAGACAA
GACTTATAAATATTAGTTTAAACATAAATCTGAAAGTGTTACTCCCTATATTTTAGTAAGACTTCTCTGTGGCCCTT
GCTCCTATAACAGTTTCTTTAAGGTTAAACTGCTGGGTTTCTGAGCTCTTCTCCATCCTACTAAATGAGAACCTCTAA
TACAAGAGGACAGTGCTTGGAATCTACATTTGTATAGAACCCAAGTGATTCTTATCAAGCAAATAATGGAAATTTCTAT
CTTAGAGAAGTCTAAACATACTAAGGCCCTTGTTTAGACTCTATGTAATTTCATTTTTCAGCCACTCTGGGCCTTGACCC
CTGAGAACCAGCAAAAGAGAACTTTCTTCAGCACTTGGCTTAGCCCTCCCTCTTTTGTGAAGGAGTTTCCCTTCACTG
GTCAGTGTTCTTGTCTTTAATGATGGTCTTGAACACCACTATGTTGTTGTACTTAAACACAGAGTTGTAACCTGTTTA
CAGGTTTATATCCATTAAACCACTTGATCTTGTGTTTCATGCTCTCTTATTCTCACCTTAGCATTTAGTTACCACTCAC
TAAATCAGTTGATCTTGAACTTTAGAAATCTTACCCTTACCTGAGGGGTCCTTGTTTACAAAGCAGATTTCTAGATGC
TCTTGCTCAGGTAATCTTATTCTGTAGTTGAAATGGGCATTTTAGGGCCAGACACTTTGGCTCATACTATAATACCAA
CACTTTGGGAAGCTAAGGCAGGAAGATCACTTGAGTCCAGGAGTTCAAGACAGGCCTGGGCAGCATAGCAAGACCCTCT
CTCTACATAAATTAGCCTGGCATGATAGCACATACCTGGCTACTCAGGAGACTGAGGTGGGAGGATTGCTTGAACCCAGC
ATTTTAAAGGCTGCAGCGGGCCACAATCATGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGATCCCATCTCTTAAAAA
AAGTGAGCATTTTAAACAAGTGTTGTAATAAGTCTACCATTTCTTAAACACACACAAACATACACACCTTGAAAATGCA
AATGAATGAATGTGTGACAGAGTGGAAGAAACAGCTTCAGAAGGGAGAATAAACATAGCATTTACAGAAGATAGTGAGA
ATACCACTGATTTGGTATGTTGGATGCTGGTTGGGAGAATTTTATGGGAAAAATACCAGTTGAGTGAAATTATGGATGT
TCTTGAGAAATAGCAGAGTTTGAATTGAATGTGGCTTTTTCATATAAAGTGGTAATTTTGGAAATTTTGGAAATAGGAGAAT
GAATAAAGATGGCATCCCAATAGTTGTCTATGGCTAGTCTTAGGAAGGGTTGGATCAGATGATCTCTAAATATCTCTT

Fig. 9.189

AACACTGAAATGTATAGTGTATGCTATAAAATATGTAGATATTGAGTTGTATTAATAAAGCCTTAGCTTGTATTTCATG
GGATAAACTTCTACTTAAATAATATTGTAATTTTCATTCATTAATGCATGTGTATAAAAGTGGATCAGTATTGTGGAAG
GTTTGCTTAAGGTGAGATACATGCAATGGTAGCCTGGAGCCAGGTAGTAGTACCAGCTCAGGAGTACTAACTGTTACGT
ATTCAATGATTTTGGAGACCTGGTTGGAAAGCACAGGCATTATTAAATTATATAAACCCGTAATTAAATAAATTATATGA
AATGCAAAGGTAATCAATACTCAAAACTCATTAGTTCCCAAGTACTTCATTATATTTTACTATTATCCATGCTCTTGAG
GTTATGTAGTCCATCGTATCTGTGTCGTGGAAATACTATATAATGAGGTGCAGCTACAAATCTCTTCCCAACTCCACAT
TCAGGGCCATCACATTGGTAGCTTGAAATAAGACATGCTGATAGTAGTTACACCATGGAAATGGACATACTGCACAAAT
CAGGCCTTTTTTTTTTCTCCTGGAGAGCCAACTGATAAATATTTACCAGCATACCAATGGCTCATGTTTAGAATAGTCCC
ATTGTTTTTGGGGTAGAAATTCATTTTGGTACATGGCCTGACTCAAAGTTTACCCCTTTAGTGTTACCTCCTGTTTCAGC
ATTGAAGCCACTTATGTCTCCTTATATGAAACAGAGAAACAAGAGGCTTTTTTTTTTTTTTAAATTCTAAGAGTGGCTGGCT
TTAGTTGTAACAGAAGAGAGCAGGTACCTTTATTGTACTTCAATTTAAACTCCTTTCAAAGGATCTGAGAACTTTTTTC
AAAAAAATTTATTCCCCCAAATCTGATAAATATGCCTCTATTTTACAATCTTTTACAACCTTTTCAAATTAATACTACAATC
TGAAAGACATTTTCAATCTTTCTTGAAAGACAGGGAGAATATTGCCATGGATGAAGAAAACAATGACTTGTCTTTCTTCA
TTGTATAAAATCATTCTTCATCACTGTGAAAATAAGAAAATGAATTTTTTTTAGTTGAGCACAATCTCGAGATCTTTCT
AACAGCCTCTGTTCTTACAGTATGGATGTTACTCTTGGTTTATAGCTCTGCAAAGGAGTCAAAGAATCTTCCTTTTATG
AGGGCCCTGAAGAAAGAAAATTGTGCTGATAAAATTACTTTATGTGCGTTTGAATGATGATATGGCCAAGCATTCTTCA
GAACTTCTGAATTTTTTCATGTATGCCTTCAAGAGTTGTAATCCTTACTTAAGCAAATGTCTGTTTAGGAGAGATTCCA
GAGATTTTCTTATTAGAGTGTTTTTAAAAATTAAGGAAGGTGTTTACCAGCTTACTAGAGGTTCCTTGTGAAAAGAGAA
ATGAGTTTTCTTCAACAGTCTTATGTCTTTGGTTCAAGCTGACATAATTAAGGTGTAAACATATCAGTATAGTTTGT
TGATTAACTTTAGATGGACATAGAATAGGTAATCAAATTCATTGGATCGAAATAAGTATTCTTACTCTGAAATGAAAC
AAAATGGAATCTTCAGAAACATGGAAACAATGACCCAAACATCAGAGAGGCATTGAAGATAAATGGGAATATCACTGGG
AATAGTGTAAATTGGAGCAGTGTTTTCCAAGCCCAATCCTCAGACCTCTCGAAAATGGAGATTGTAACACTAGATTGTGG
GCAGTCTATACCAAACCAACAGTACTGAATCAGAGTGGGTATGGGAATGGCTAGAGCAATTTGCAATTTTACAGCATAT
TCAGAAGATTTTTATGTACACTGAAGTTGAGACGTGCTGATTTAGCAAAGGTAAGTACAATTTGCCTCAATTCCTCA
TCTGCAAAATGGGGATTATAAAAGTAGGGTTGTTGGAGGACTGAATGAGTACATATATGTATAGTGTTTAGGACAATGT
CTGCTGTTTTACAGTAAGTGCTCTATTGTTGGCTCTTGCCACTATTATTGTTTTGTCAAGGGCTTGTCTGTGCTAGCTG
GAATGGAACTTAAATGTTTTCTAATTATTGCAACTGTCAGATTTCTTAAATATCGTAATGAAGCCAGCACAGTGAAGAG
CTGTCCTCAGTTTAAATGTAACATTGGATCTATCCTAATAATTTTTCTTAGTTTCTTATTGCAATTTGTAACATTTTTATT
TAAATGTTTTATTTTTCCACAGGGTTACATTTATTTTATTACTTCAGAAATATTAGATGTCAATCTGAAATGTGGCTG
TGCCTTTATCTTGCATGTGGAATTCAATTTTTTATATTCTTTACAAACAGTATTTATAAGAAATAAAGATAAGGTTACA
GTGAGTTGTAGTATGAGATGGTAATTTACATGGGGTGCTCCATGTGCTTAGCTTTTCTCTGAAGTTATGCATCCCCACA
GCAAGGGAAATATTTGCATTTCTGAGAGTGAGAATTTATGTTTCATCCTCTACTAGGAATGGTGGCAGCTTTTCCAGGTC
AAGGCCCTGCAGATGCCTGATGGTCATGGTAAATGAAGGCTGGATGCAGGGAAGGCAGCAAAATGAGAAAATCTCCTGG
GATCAATTAATGGGAGTCATCTGAGAGAGAATAAAGGCAGGAGAACATTTTCTTCTAGTCAGGAATTCGCATCAGTTTTT
GCCTGGTAATGGGTTTCATGAAAGCCAAGATGAAAGGGTTTTATCCCTAAGGAAAAAGGGCTCTCCTCACATCCTCTT
TCTGTGCTCTTTTATCAATGACTAATAATAACATGCTTGCATCCATGACAATCTTTCAGAAAACCTGATGCAAAAACAAG
TCCAAAATTAGAGTGACAGCTGTGCTGATTTCTGGAGTTTCTGGTGTGTGCTTAGTAGGCACACAGTTTGAGTTACA
TGGCTTCATTATATGATTACACACTACTTTGTTGCCTCATTGGTGGTTAAGGTGATTTTAAACTTTTCAGTACCCAAAA
GCAGTGTCTGGAACATAACCAATGAAGACGGAGAATTGTTTCAGTCCCAGAGATATCCCAAGCAAATACAGCAGTGAAC
TCAATGGATATTTGTGAAATGAATAAATAAATGAATGAATGAGTACTACTGTGAATATTCTTGTGATGTTTGTCTGTTTC
TCATTTGGAGAGTTTCTTTGAGGTCCGTGACCAGAGGTTGAATTGCTGGATCTTAAAGTAAGAGGATTACCAACTTCCC
TATACGATGCTCAATTCCTCTCCACAATGGCTATACTAGTTTACTGTCTGGCAATCGGTGTATGAGGGTTACCTGTTTA
TCACATTGTTGTAACATTTGATTTTGTGATGTCAGGTTCTAACTGAGGTCCAAGGGGAGTTGGTGGGCAAGTGGCGG
GTAGCTGGAAAAACACGTGAGGAACGTGAGACAGTTTCAACATGGATTTACTCTCTCTCTGGCCATGAGCCATGAGCGC
AAGATGTATGTACAGCGTCAGCAGCGTAGTTGTACCTTTTACAGACAATAGTGGCTCCAAGCCAAGCACAGCTTACAT
GGGTGATCACCTAATGTGCCTCACGTGGCGTGGTTACATAATGTGCAGAGTTGTGAGCCTGTGCTCCAAACTCACTGAG
TCATGCTGGACCGGATGTCTGCTTCGGCCTATTTTTGAGAGCAGCACATCCATTTTCTTACACTCCACCCTCAATGCC
AAAGGAGACACAGGCCTTGGACACACAGGTCTAAGACACAGGCCTTATACGTACACTCTGGGGACAAAGGCCCTGGACA
CACAGGTCTGACACATAGGCCTTAAATTTCTACTCCCTAGGCTGAAGGAGTCTTTTAAATGGAGAAACATGCCCCACAGGG
TGGAACCCAGACCCAGAGGCCACAGCAGTAATAAAGGAGCAACAACCTCCAGGTTATGACGGGCAACACCCCATGAT
GATGTTACCCGAATTTACTTTATGCATTAAGCCAGGTTTTTATTTCCCTACCTTTAGGGGCATTGGGGCATGCAACAAC
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TGTCCATACCCGGTATGAAACCATATGGAGTCAGTACTAGGCTGGACTGTGAGCAGTCCAGGCAGTAGCAGCACCCAG
CTAGACCCATCTGTGTACCGTGCCCCATCGGGCATAGGGGGGGATGCCCTTCTTAAATGGTGAAGGCTCAGCATGTAG
GGGTGCTTCAGGCCCCATGTCCTTAGGCCCCATGGCCTTATCTTGCAATTAGGACTGCTGGTCTCAAGACCTCTTGCAAC
TCTGCTGCTAAGGGACTTGTACTCAGCGTACTCTGTTGCTTCAAGTAGGTGCCCACTTTGCTAAAGTGGATGTCTGCA
CTGTCCCAGTCCGGGGGGTGTATCAATGAACACATCCATCCTGCTATCAGGTAACCTCGTCTACACGACAACCTGTAGC
CTGTCTGCCATGCTCTCATGAGCCCGAAGGGCAGCATATGCAGTTACTAACTGCTTCTCTATCAGTGAATACTGGAGC
TCAGCTTCTTTCCATAGTTGGGACCAAAAGCCTACTGGTGTCTCAAGCACTCCATGCTCTGCCACAGGCCCCAGCCAA

Fig. 9.190

Fig. 9.191

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CATAGTTCTGGTTCGTTGTWTAATTGTTTATATCCTTGTAGCATATTCAGATGCACATAGTCACCATGTCTCACCCAGA
ATCAGAATCTCTCAAGGAAAGAAAGCATGTCCTCTCTTTCCCTAATAATCCCCAAAGCTCTCAATACCATGCTGTGCCC
ACAGAAGGGAATCAGTTAAATCTGCTGCAATTGATTGGAACTTCTTTTCCCCCAGATTTTCCTTAGGTGGTCTCCTGC
AGAGTTCTGTCTAAAGTAATGGAGACTTGGGATTTGTATTCTCATTATGCTAATGGTTATTACTTCTTTTATTTTTGAA
AACTGGTTGTAGGATCTAAGCTAACCATGCTATTTTCTGCATACCACCCAGCGATTCTCATTTAGCAACTGCCTTCAA
TCGTCTACTCCCTTTGGTCTCCTCCTCCCTCAGTAGGAGAAGGATGAGAGGAAGTTGGAATATTTACACTGAGATGAG
ATCCTTGAGCACTTGAGCCCTGAGGCTGCCTCTGGACTAACGTTTCTGTTCAGCTGTCCCCAGGTTTTTCAACAAGAGC
TGTCAGAAACAAGTTTGTGACTAAAGAGCAGCTTCATCTGTAACCTTCTGTTTCTTGCTACTCCCTGCTGTGTTTCTGGC
TATCTGGAAGCTTCTCAGCAGGGTAGCAAGTTGTTGGCGCCACTAACACCTTTCTCTTTCAAATTCATATTCTTCCTT
GGGAGGTGAGTTTGCATTTTCAAGAAATTTCTATTTTAGTGAAACCACCCAGGAATCAGTGTGAGTTGGCTTTTCACTGTG
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GTTCTTAGGAATTTTATGCTTAGCTGAAAGTATTAAACACTTTTCTTATTTCTCAACATGAATTTCTTTTCCCGGGG
CTGCAGGGGAAAGGGCTCTGATCCTGGTTTCACTCTTAGCTTGATCATTTTAGTCGAGTTTCTTTACTTCCATGCAACT
CTGTTTCTCTCAAGTATAAACTATTAGTTGAGTGGGTAGAAGAAGTAGATGATCTTTAAAGCACTTTTAAAGTCTAAAA
TGTTATGATTCTGCTATCAAATTCACAGCTTCTCAACATCAGATAAGGGTTGGCATGAGTGTGCTGAAAATATATTTG
GGTTGGAGGAAGGGATATCCTCGATTAAAAAAGCCAGATGAGAAATTAAGACTTTTTCAGCCAATGCTATT
GACATAAACTCAGGGTGACTTTTACGTCCTAAGTAGAAGTGGAACGTTTCAATTAAGCTTAGGTAATACCTCTTTA
TCAAATTAATAATTTATGTTCAAATTAATCATCATGAATTGACATAGCTAATTTGACCCTTCATGGTTATCAGAGTCT
TTTTGGTGAATATCATAGTCTACATTAATTGAAAGAGGATAAGGTACATAGGTGTCTCTCCCACAACCTAAGCCATCAT
GTCTAATGTAGTAATGAAAGCAGATTTTGGACTGAATTTTGTGACTGGCAAGCTTTTGTGTTACCAAGGAGCCACAAC
TCATAAAGCCTATTCTCATAAGCTCCATAGACTCCCAGCTTTCTGTCCCTTAGACATTGTGATAAGCGTCTTACATGGA
AGATCATTTTTTATTATGTCTGTTTGAATTTCTCTCTTTCTTCTTATTAGTCTAGCTAGTAGTCTATCTATTAATTT
TTTCAAAAATCAGCTCCTGTATTCTTGTATTTTCAAAGGGCTTTTCTATGTCTATCTCCTTCAGTTCTGTTCTGGATC
TTGTCTTCTACTAGCTGTGGGGTTTGTGTTGAGAGGACATGAACAGACACTTCTCAAAGAAGACATTTCATGCAGCCAAC
AAACATGAAAAAAGCTCAACATCACTGATCATTGAGAAATGCAAATCAAACCACAATGAGATACCATCTCACACC
AGTCAGAATGGCCATTATTAAATAGTCAAGAAACAACAGATGCTGGTGAGGTTGTGGAGAAGTAGGAACGCTTTTACAC
TGCTGGTGGGAATGTAAATTAGTTCAACCATTATGGAAGACTGTATGGCGATTCTCAAAGATCTAGAACCAGAAATAG
GATTTGACCCAACAATCACATTACTGAGTATATACTCAAAGGAATATAAATCATTTCTGTTACAAAGATACATACACAGG
TATGTTTATTACAGCACTATTACAAATAGCAAGGCATGGAATCAACCCAAATGCCCATCAATGATAGACTGGTTGAAGA
AAATGTGGTACATACACACCATGGGATACCATGCAGCCATAAAAAGGAATGAGATAATGTCCTTTGCAGGGACGTGGAT
GAATGTTCTCACTTATAAATGGGAGCTGAACAATGAGATCACATGGACACAGGGAGGGGAACAACACACTGGGGCCT
GTCAGGGAGTGGGGTTTAGTGAGGGAAAGTATTAGGAAAAGTAGCTAATGCATGCTGGTCTTAATACCTAGGTGATGGG
TTGATAGGTTTCAGCAACCACTGTGGCACACATCTACCTATGTAACAAACCTGTGCATCCCGTGCATGTACCCCGAACC
AAAAACAAAAAATAAAAAACCTAAAAAATCTTATTTCAAACAGGCTATGGGCTAGATGTGGCCTGAGGGCCATA
GTTTGTAAAGCCTGGGACTAGACCAACATATGCATGTGACCATCTTGTGGATGAGTTTCCACAAGACTGAGAGCTCCA
TAAAGGCAGAGACTGTATTTTCTCATTTTCTTATTTCCCTCAGTGCCAAGAAAAACAGTAATTCAATAAATGTTTGTGTC
ATTAAAGAATTTATTACGAAGAGTTATAAAAATCAATCCACTGTTTTTCCATTTGCTATCTCTGAACATGTACTATGT
CATTTTTACTTGAAGATATTTATATTGAAATCTCTGGTTAGTGATGTCTCACTTCTGTGTACAGCTAAGAAGACGTACA
GTTTTTAAATAATGTTACTAAGTAAACAAAACCTATTTTATCAAGAAATATAAAGAAATTTCTTTAAACAAGGTTGAG
AGGTTACTTTCATGCATTAAGTTTCTGCTATCTTGTGCTGATGTTCTGCTATAAAACCTTACAGACACTATGTCTGTATTAC
TGGTTTCAATTTGAAATTTTGCCTTATGATTAATCTAATTTGGTTTCAATGTTCCATAATAAAGCAATTGAGCATCACT
GATGTGSATACCATTTTCTTTAGGTTTAAAGAAAACCACTCTAATATTTTTCAGGGCCATTATTTTATCAAATATGTA
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TATGTTCTCATTTCTTGCTTAGAAGAGAGATATTTCTAGTTTTCTTATTTGCTAACCTTATAATTTGAGCACAGAATGTC
AGATTTGTTTTCAGCCCTTCGATTTAATCACATCATGTCTGATATCCTAGCATTTAATTACCCAAATTCCCAGTGTTGTT
CAAGAAGGTAGGTCTGTGGTGGGTTAGCCTTAAGCTAGTGGCTGAAAAGGTCTTGATTTGCCCCACATCTAACCTAGT
CACTTCTGAGCTGAACTCCCATATCACCTGGGGACAAGCGACCTGAAATCCTCACTGGCCACAGCTTCTTCTGAGG
GAAGCTGCAGGAAAAAGAACAAAAATGAGCTTTTGTGAAAGGATTAACCTGGGTGAAGAACATCCTCCATAACCAAATA
AATGGAACCTAGTAACCCACATGGCCAATCCCCAGGGCACTAGTGTCTTTGGGGATATCAGGGTGTGTATAGAATTTG
TTTTTGTCAATTTGAAAACATAAAAWAATCTCTGAAAGAGTACTAGTGGTCAAGGTAAGTCTAGAGATGGACCAA
CAATCATATTAAACCTAATTTAATCTCCAGAATTTATATTTTGGAGATTGGTAGCACTACTATCTTCATTTTGTACTTTC
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CCACTTTGCATAGTTGATCCTACAAGAGGATAAGTAGAGATCAAAGTAAGAAGTGATCAATGGATTACTATGTCAAGCA
TGATTCCCTCAAACCTATTGGGAAGAATAAATACCATAACATGTATAGGGTGCTGTGTACAATTCCTGGCACAAAGTTAA
GTACATGATAAAATGGAACCTATTAAATATTAGTAACCCAGATAGGTTCTGGGCACATTAGCAGGAAGGTCTTCTTAATT
TAAGTAAGGTGCTAAAAGCATCTAATTTTATTAGGATTGTTATGAAACAAAACCTAAATGTGCAAGAAATATATTTAC
CGTAGCCTGGGGAAAATTGTCATGGTTTCTCTTTTCAAAGAATTCAATTTGGTAAGAAACAGCCTTCAATGGTCCATGA
AAAGTGAGAAGTGAGGGAATAAGAGGCTCTTTCTATGTTTGCATTTTCAAGGCTTGCCTTCTTACTTTTAAATTTATCT
TTACAATTGCCTTCTGCTGGGTTTGTGTCTTCTCAGTTTAGGGAAGGTGACAAGAAAAGTCTGTTTGTGTGTAATT

Fig. 9.192

GACATGAAAAATGGTAGGCTTTTAACTCAAGGAGGCTCCAGACAACAGTTTGGTTATTAAGGAAGGGGAGGACCTAGGG
GAACAATTGTTTCTGGCCTCTCTTCTTCTCAGGCCTGGAACAATCTTTCATTTACTATATTCTTGCCACAAGAGCACAG
TCAATATTAAATTCCAAAAAAGAAAAATCCCTCACAGAAACAGCTGTGTTCAATATACAGCCAGTTTACTCTGGCGAA
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AAGGGAAGGCTGGTTTAAACAGCTTTGACTAATATTGGTAGTTGGAGTATTACAGTATTACAGAGTTTGTAGAGAACTAA
AATAAACCTGATGGAAACACACACATATTTCAAAGGAAACATTTTAAGGTGTTCTGGTTCCATTTTTCATTTTGTAT
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AAATCTGTCCATGGCAGAAAGATTAGTGATTGAGATATGGTGTTCATCATTGTAGATGATGGTGTCCCATTAAAGCATGT
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GCTGGAGGTCATGTGTTACATTTCTGTTGTGTAACATGGCCTCCCTCTCTTTCTGTCTCACTTTCTCACTTTCTCC
CTCTTTCTTTTACTCTCTCTGCTCTACCTCTGACTCCTTCTCCTTCTTCCAGCTTTGCATCAGAGCCATGTAAGTGA
CATTTTCTTGGTTCCCTCAGGACCTGTCATAATAGCAGGCAAATATTAGATGATGATTCAATGTTTAATGAATTTTTCT
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AATCTCTCTCTAGTAGACCACAACAAGTCAATCTGTGCTCTTTGATTACAGTTGTTCAACCAGTTACTAATCTAACTAA
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GATTATTCTAAAGTTATTTTTTCCAAAATAACTTTATTACTCTTGATATCATAACCATATTACAAAACATTTCTGGTAA
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GTTTTCTGAGTAATTCCAAACTGTGTAAGAGATTATGTTCCCTTTGCATATTGGCTGCTAAGAAGCTCACTTTTTCTACT
TATGATACTTGGATGCATTTTGGCTTTTTTGGGTTTTTGTGTTTTTGGTCTAATTTTTTTTTTTTTTTTTTTTTTTGA
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CAGGTGATCCTCCACCTCAGCCTCTCAAGTAGCTGGAATTACAGGCATGTGCCACAACACCCAGCTAATTTTTGTATT
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TCCCAAAGTGCGGGCATTACAGGTATTAGAGGTAGGAGCTACTGCATCTGGCCATGGGAAACATTTTGAATGATACTT
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AGCACTTTGGGAGGCGYAGGCAGGCAGATCACAAGGTCAGGAGATCGAGACCATCCTGGCTAATATGGTGTGAAACCCC
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ACAAGATCAGATTCTAGGGGCTGAATGGAGAATAAATATGTAAGTGGCAAGATTGAAGGCTGGGAATGTAAAAGGTGGC
TCCTGCTGTGGCTTGGGCAAAACATCACATAGGCTTGAGCTAGCTTTGTGATTCTACGGGATTGAGAGGAGAATGCTAA
TTTCATAAATGTTTGTAGTGGATATGATTGGCATGTGAWCTGAATGCAACTGGAGAAGGGGACCAGTCATTTACTGAGTWG
ATATTGGTAATTGATAAATAAGACCAAAGAAGGTGAGGTTTGGGGATAGGCAGTGTAATTTTGGGAAGTAATAAGGGGA
TTGAGGTTCTGTCCATGGCATTGGTGGGGAATCTAGCATGTAAATACATGTAAATAGATAAGTAATATGCCATATACA
TTTATATAAGATATCTATACACTTATTCTCCTTCTTTGTTTATTATTAGGTGTGAAATCTACGTGATTTTTTTCTCCC

Fig. 9.193

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ATATTATTTTTGGTATTCAATAAATGTGAATCATGAAGAGATAGCTGTTTTATTTAATCTCTGCAAAGTCACATTCTTA
CCCATTTTTGATTGTAATAAAAGGCCTATGTCGTTATTTAATTTTTTAACCTTTGTGGCATGATTTATAGAAGGAAAATA
AAAGTGTCAATAGGTATGATAGGCCAAAGTTGGGTTGTTGGTAAAATAATAATACCACAAAATGTTTTTCTATTTAGCCA
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CAGTAATGGTTATGCAATTTCAATTGTGCAGGGTATTAAAATTTGTGACCAGGGATCCCAGGAGACCAGCTATTAGATTT
TCAATGCACTATTAGATTAAATMTATCTACTCAAACCTAAAGGGATCCTGCCTGAGGCTGTCTGATCAATAGTCTATCAT
TCCTGTGCAAATGAAGCTATTAAGATTCTCTTAGGAGGTAGACTATCTAAATTGGATCTAACTAGTAGATGGTGACTGA
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AGGTTGAAAAAGAGGTGTAAAATAGTCCCTATTTTATATTATTGGTTCTTAAGTACTTATCAGAAAAATGACAAAAGTC
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TGCAGTCACCATTACCTCCACCACCATCATAATTATATATAGCTACCACCTTCTTGACACTCAGGTCTGCTTGTGTTG
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ACAGGAAAAGCTGATTCTATAATTTAAAAATGATTTGGGGATCCAATAAGTCATGATTCTATTTTATATAATTTTGGA
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TTTATTTTTATGGGTCTGAATCTTCACTAGACTAACTGCTTGAGAGAGGAGCATGTATAATTCATTTTCATGTTTCAAA
TGCTGAAACTGATGTCTATAGAGCTTCAGTATTTGTCCACTGCATAAAATCAGTAAATAAAGGCAGGAAAGTTTGTTAC
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GCCCTGACCTGAATCAACTCTGAGCCCCCTACCTAGGCTGTTGTTTCTTAGCAAGTACATAATAACAACCTCATCAGAATA
GCAGAGATCTTGGTAACAGGTTGCCTGACACTTGACTGATATGGGAAAATCCTGAACAGGTTTGGACTGGAATCTTGGC
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AACTTTGTTTCAATTGGTTTGCTTTGAAGAGTAAATGAGATGGTAAGTGTTGGTGTATGTAGTGCAGGAACATAACAATTC
TTGATAGATATTCATTGAATACCTRTAGTACATAGAACATAGAAAGTTCCTGTTATTCCACTCCCACCCCCACACTT
GCCCTTCAGGTCTCAGGTTCTCCTTCTGTTACCCCCAGAAAGATTAAGCCCCCTTTGGCCCTCTATGTGCCTCTTTTAT
AGCATTTAACACAATTAATAATTAATTGATTATGTAACCTTTTTTATTTAATAGCTGTCTCTTGTAAGATATAAGCACATA
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CTTATTCAACTTTCTCATCCTCAAGATTAGCAAAACTACTCTCTTTCTATTCCCAAATATCTTCTCTCAGTTTCATTAC
CATGATCAGAAAGTACTCTTCTTGAATTTATTTTCTAGACTAGATCTATTTTGGGGTTTTCTTACCAGTGATTGTAAC
TATTGATGTGGTCTTTTTTCTGTTTTTTACTTCTCATATGTGTATCTCTTCCATTAGTCTTTCATTCTTCAACAGAT
ACTTTCTGTTGTGACAGCCACTCTTCTAAGCACTATGGAAGGATTTAAAGCTGTGTGAAGCACAGTTATTCCGGTCAAG
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AACCAAGAGCTTCAGTATTTTCTGCAAGTTTTTTGCTCTATTGTATTCTCAACTTTTACTGGTTTTCTCATCAAATAAAT
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CAGGATGTGTAGTCAATTGATGTATCCCTAGAATAAGTGCCCAATTATAGTAGTCATTTGATGGACATTTATTAAATGAA
ATTATGAATGAAGTAGGGATTGGCAGTTATACACTAAGACTCTGGAAAAATGGCCTGAGTTGAATCCTGGCTTTACCAT
TCTTGGCTATATAATCTCATATTATTTACTTTTCAAGCTTTTCAATTTCCAGATGCCCTTACCTTCTAGGGTTGTTGGGAG
GATGGAATGCATAATACATGTAAACCAGCAAGTTCATAGTAACAAAAGTTGACTTTTTTAAAAAGTTAACTTACTCTTCT
TTCCTATATATGTGTAAATCATATTTTATTTTCTCATTTTAAAAAGAAGACAATAACTAAAGGTATTAGCACATAATGA
TTCAAAACATAATATTTCCACATGTGAGTTGAATGACTTTTGAGCCAGATAATATGAGTTGAAATCAGTTTAAACAATA
TTAAATCAGCCAAGAGCAGTGTCTCATGCCTGTAATCTCAGCACTTTGGGAGGCTGAGGCAGGTGGATCACTTGAGCTC
AGGAGTTTGAGACCAGCCTGAGCAACGTGGTGAAACCCCATCTCTACAAAAAATAACAACAACAAAAAATTAGCCAGGTG
TGGTGGCTCATGTCTGTAGTCCAGCTACTTGGGGGGCTGAGGCAGGAGGATCACTTGAGCCCAGGAGGTCAAGGCTGC

Fig. 9.194

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AGCGAACTGTGTTTGTACCACTGCACTTCAGCCTGGACGACACAGACCATGTCTCAAAAAGAAAAAATAGAATAAAA
TCAAATACACACCATTTTTGAATATGTCACCAGTCTGTGTAGTTTCATCTTGAAAGGACTTCAAGGTCCAATATGCATCA
TGCAGAGAGTTGCTAGGGCCCAGACAAGAGTGAAGTGGCCTGAGTAGTTAACTATCAGATAACTAGTGAAA
CAATTCGGCTATTTCAACAAACATGTTGCATACATAACGTGTTATGCACAGAGATGACATTTGGTATAATATTATGGCA
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AATGAGGGGTGGGTGAGTAGGGTAGCAGTTGGCAGAAGACTCACACTGGAATAAATGCTCCTTGACTCGATATTTTTTTT
TTTTTCAAAAATGCAGAAGCATTATTGAAGCCCGGATGTTTGGGTCTATGATAAATAAAAGATATGGATTTGAGCTGC
TCATTAAATTTTGAAAGAAAAGATCCTAAAAGGTTAGAGACCATGGAGATTTCAAAGTGGCTGACCTTGATTAGATATA
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GTGCCTCCTTGGTGGAGAGCTGACCTATGAGTAGTTACTGTGTGAATTAATGAACATCCTTCAGCAAAAGTTATTAATA
GTAATGTTTGGTAAAAGTCCTTTAGAAGTAGACTGTTATGTGTGTTACTAGTTATAATCAATTAATAACCTGTGATTTG
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CAAGACATTGGCTTTGTTCTGAAGCAGCTCCACGCTCTTCCAGAAATCTCTATGCGGGACTCTGAATGTGGTCAAGAA
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CACACAACAGCTGAAACATCTTTTCTTCATTTCTTTTAATTCCTGTAGCATTTGATGTCTCCACCGTGTAATTTACATT
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GAATAAATACACTGAGTTATTGGGAGTTTGTGAAGGAAGTGACTAGAATTTCAATAAAATAATAAAGTTTGTGTTTGT
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GTGTCACAGACTTTGTACTGGAAAAATGTCTGCCTTAATCTAGTGAATAAATATGACTGTTCTTTGAAGTCCTTTTTTT
CCTCCCCGAGTATAAATTCAAATGCTATTTCATTTTCTGAGTTGCCTGTATTTCTTTAGCCCTTAAGGCATCAACCTTTG
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CAGAACCTTATTATCATCACAAGGAAGTAGATTAAAAAATACTTTTTCAGTCATTCGTATTCAACAAGTACACTCCATCA
AATCTTGCCTAACTTTTTTTTGCAAATAACTTTGCTCCTTGGATCCTCTCCAGGTCTTTATCAAAATGGAACCACATAC
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AAGTTCCATTTGTTTCACTAAAATTYGCAAATAGTTGTGATTTTCTTTGCAGATCTGTCCAGTTTGATCTTGAAACAAA
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CTTAAATTAGAGCTCAGCCAACATAGAATCTAGTTCAGCAATACTCTACAACATGAGATAACCATACTGATGTTTGATA
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TACGTGAGAGTAATATGAGGTGATGGGGTTTACTCTTAAAGAGATTACTAATAATGTTTATTTGGAAAAGATGAAGAT
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TTTTTTTTTTTATTATACTTTAAGTTCTAGGGTACGTGAGCACAAAGTGAAGGTTCTGTTACATATGTATACATGTGCCAT
GTTGGTGTGCTGCACCCGTTAACTTGTCAATTACATTACGTATATCTACTAATGCTATCCCTCCCCCTCCCCCACTC
CATGACAGACCCCGGTGTGTGGTGTTCACCCCTGTGTCCAAGTGTCTCATTGTTCAATTCACCATATGATTGAGA
ACATGCAGTGTGTTGGTTTTCTGTCTTGCAATAGTTTGCTCAGAATGATGGTTTCCAGCTTCATCCATGTCCCTACAAA
GGACATGAACCTCATCATTTTTTATGGCTGCATAGTATTCATGGTGTATATGTGCCACATTTTCTTAATCCAGTCTATC
ATTGATGGACATTTGGGTGGTTCCAAGTCTTTGCTATTGTGAATAGTGCTGCAATAAACATACATGTGCATGTGTCTT
ATAGCAGCATGATTTATAATCCTTTGGGTATATACCCAGTAATGGGATGGCTGGGTCAAATAGTATTTCTAGTTCAAGA
TCCTTGAGGAATCRCCACACTGTCTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAACAGTGTAACCTGTTCCCTA
TTTCTCCACATCCTCTCTAGCACCTGTTGTTTCTTACTTTTTTAATGATTGCCATTTTAACTGGTGTGAGATGATATCT
CATTTGTGGTTTTGATTTGCATTTCTCTGATGGCCAAGTGATGATGAGCATTTTTTTCATGTGTCTGTTGGCTGCATAAAT
GTCTTCTTTTGAGAAGTGTCTGTTTCATATCCTTTGCCCACTTTTGTATGTGGTTGTTTGTATTTTTTCTTGTAATTTGT

Fig. 9.195

TTAAGTTATTTGTAGATTCTGTGTATTAGCTCTTTCTCAGATGGGTAGATTATAAAAATTTTCTCCCATTTCTGTAGGTT
GCCTGTTCACTCCAATGGTAGTTTCTTCTGCTGTGCAGAAGCTCTTTAGTTTAAATTAGATCTCATTGTCAATTTTGGC
TTTTGTTGCCATTGCTTTTGGTGTTTTAGTCATGAAGTCCTTGCTGTGCCTATGTCCTGAATGGTATTGCCTAGGTTT
TCTTGTAGGGTTTTTATGGTTTTAGGTCTAACATTTAAGTTTTTAATCCATCTTGAATTAATTTTTGTATAAGGTGTAC
GGAAGAGATCCAGTTTCAGCTTTCTACATGTGGCTAGCCAGTATTTCCAGCACAATTTATTAAATAGGGAATCCTTTCC
CCATTTCTTRTTGTTGTTCAGGTTTGTCAAAGATCAGATGGTTGAAGATGTGTAGTATTATTTCTGAGGGCTCTATTCTG
TTCCATTAGTCTATATCTCTGTTTTGGTACCAGTACCATGCTGTTTTGGTTACTGTAGCCTTGTAAGTATAGTTTGAAGT
CAGGCAGCGTGATGCCTCCAGCTTTGTTCTTTTGGCTTAGGATTGTCTTGGCAATGCGGGCTCTTTTTTGGTTCCACAT
AACTTTAAAGTAGTTTTTTTTCCAATTCTGTGAAGAAAGTCATTGGTAGCTTGATGGGGATGGCACTGAATCTATAAAT
TACCTTGGGCAGTGTGGCCATTTTCATGATATTGATTCTTCTATCTATGAGCATGGAATGTTCTTCCATTTGTTTGTG
TCCTCTTTTATTTCTTTGAGCAGTGGTTTGTAAATCTCCTTGAAGAGGTCTTCCACATCCCTTGTAAGTTGGATTCTTA
GGTGTTTTATTCTCTTTGAAGCAATTGTGAATTGGAGTTCACCTCKGATTTGGCTGTTTGTCTGTTATTGGTGTATAGG
AATGCTTGTGATTTTTGTCACATTGATTTTTGTATCCTGAGACTTTGCTGAAGTTGCTTATCAGATTAAGGAGATTTTGGG
CTGAGACGATGGGGTTTTCTAAATATACAATCATGTCTGCAACAGGGAGAATTTGACTTCCTCTTTTCTTAATTG
AATACCCTTTATTTCTTTCTCCTGCCTGATTGCCCTGGCCAGAACTTCCAATACTATGTTGAATAGGAGTGGTGAGAAA
GGGCATCCCTGTCTTGTGCCAGGTTTCAAAGGGAATGCTTCTAGCTTTTGCCCATTCAGTATGATATTGGCTGTGGGT
TGTCATAAATAGCTCTTATTATTTTGAGATACATCCCATCAATRCCTAGTTTATTGAGAGTTTTTAGCATGAAGGGTTG
TTGAATTTTGTCAAASGCCTTTTCTGCATCTATTGAGAGAATCATGTGGCTTTTGTCTTTGGTTCTGTTTATATGCTGG
ATTACATTTATTGATTTGCATATGTTGAACCAGCCTTGCTATCCAGGGATGAAGCCCACTTGATCATGGTGGATAAGCT
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TTGTTGATCTTTTCAAATAACCAGCTCCTGGATTCAATTGATTTTTTGAAGGGTTTTTTTTGTGTCTCTATCTCCTTCAGT
TCTGCTCTGATCTTAGTTWTTTCTTGCTTCTGTTAGCTTTTGAATGCGTTTGTCTTCTCTAGTTCCTTTTAATT
GTGCTGTTAGGGTGTCAATTTTAGATCTTTCTGCTTTCTCTTGTGGGAATTTAGTGCTATAAATTTCCCTCTACACAC
TGCTTTAAATATGTCCCAGAGATTCTGGTATGTTGTGTCTTTGTTCTCATTTGGTTTCAAAGAACATCTTTATTTCTGCC
TTCATTTCTGTTATGTACCCAGTAGTCATTACAGGAGCAGGTTGTTTCAAGTTTCCATGTAGTTGAGCGGTTTTGAGTGAGTT
TCTTAATCCTGAATTCTAGTTTGATTGCACTGTGGTCTGAGAGACAGACTGTTGTAATTTCTGTTCTTTAACATTTGCT
GAGGAGTGCTTTACTTCCAACATATAGGTCAATTTTGAATAAGTGCAATGTGATGTTGAGAAGAATGTATATTCTGTT
GATTTGTGGTGGAGAGTTCTGTAGATGTCTATTAGGTCCACTTGGTGCAGAGCTGAGTTCAATTCCTGGATAACCTTGT
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CTGTTTTATCAGAGACTAGGATTGCAACCCACCTTTTTTTTTGTTTTCCATGTGCTTGGTAGATCTTCCCTCCTCCCTT
TATTTTGAGCCTGTGTGTGTCTCTGCACGTGAGATGGGTCTCCTGAATACAGCACAGTGGTGGGTCTTGACTCTTTATC
CAATTTGCCAGTTAGCATCTTTTAAATTGGAGCATTTAGCCCATTTACATTTAAGGTTAATATTGTTATGTGTGAATTTG
ATCCTGTCATTATGATGTTAGCTGGTTATTTTGTCTCATTAGTTGATGCAGTCTCTTCTTAGCATCGATGGTCTTTACAA
TTTGGCATGTTTTTGCAGTGGTTGGTACTGGTTGTTTCTTCCATGTTTAGCACTTCTTTCAGGAGCTCTTCTAGGCCT
GGTGGTGACAAAATCTCTCAGCATTTGCTTGTCTGTAAAGGATTTTATTTCTCCTTCACTTATGAAACTTAGTTTGGCT
GGATATGAAATTTCTGGGTTGAAAATTTCTTAAGAATGTTGAATATTGGCCCCACTCTCTTCTGGTTTGTAGAGTTTCT
GCCGAGGGATCAGCTGTTAGTCTGATGGGCTTCCCTTTGTGGGTAACCYGACCTTTCTCTCTGGCTGCCCTTAACATTT
TTTCTTTCATTTCAACTTTAGTGAATCTGAAAATTATGTGTCTTGAGATTGCTCTTCTCGAGGAGTATCTTTGTGGCAT
TCTACTAGAGGGTAATTTTCTTTTAGGGTGTATAATTATTTTAAAAATTGATTCATTGGAATATGTTATCAATGTCAA
AACATGAGATAACTTGATGATTATCTCTTTATGAATTTGAGGTGCTGAGTATTTCTTGTTTTTCAATTACAGCATTTCTT
TCAAATAGCCAGTCTCATGTTTCTATTTCTCTCTTAGCTGCTTCTTTCTTTAATATTTTGTGTGCTAATGAGAACCCTTA
CCAAAATGAGTAACCAAAGGGCTGAAGAAAGTTGAACTGTAAATGGTGTGTTGTTAATTGCTTCTTCAATTACAGAAGG
ATTTTATATGTTGAGTAACATGCCATTTGAAGATTTGACAATTCATCCTCTCCTTGGTAACTCCCCAGTCATTTGGTC
TAGATTGATCCATCATCTTTCTCACTTAAGAGCATTTAGTTCTCACCTTCAAATAATCAGCAGATTAAAATAGTGGTAC
ATAGTCATCTGTAGGAAGAAAGAAATAAGCTGTACAGTAGAGTGCTCTGTGGGTAAATATTTGTTGTGTTGAGATATAA
CCTGTGTATATTTATTAAATATCAGTGACATTTCTTTGCTTATTCTTTTCAATTCACCTCAACAGATATTATTGAACAGCTG
CTGTGTACTAGGCATTGTTCATTTTCCCAAATTTCTAAATTTGTGATCATTCTATTTTTTAAATGTATTATTTAACAAA
GTACTGGTAATGATATATATCTTTATTTCCAGTGGGTTTTTGGCAATATAGCATAAAGATCATGAAAGTGGGCTCTAGA
ATCTGAGAGCCTGTGAAAACCTTCAGCTCCAACACTTTCTAGTTGTGTGAGCGACTTTAGGCAAGTTAATGAACTTCCCT
AAGCTCATTTACTTTCATCTTTACAATAGGTATAATAAGTACTTACCTCATGTCCACTATCTAAAGAGTAAATGGGAA
AATATGCAAAGCATATTGCACTGGGCTTGGCAGGCAAGTAATCCCTCAATAGATGTTAAATTTGCTGCTGTTATTAAAT
CATAGTTCTTGCTTAGTTGATTGTTATAATCTTTTTTCAAGTATTTATTGAACAAATACCTATGAAGTGTACTCAGAGTAA
ACTTGGACAGTGAGAGACACCACCATCTGGCCTGTTGTGTCTATAAACAAGCCTTAAACTGGGTCTTGGCAATGGTG

Fig. 9.196

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AGGGGAACAGATGCTCTAACTGCCTTCTAGAATGTGTTTTTCACTATGTGTCTAAGCTGCGATGAATGTCACCTAATGTC
ATCTCTCTTTCTATTTTAGGTAAGTACATCACCATTTTTTGAACTCTGTGCAGATGGGTAAGTCTTGCTGCTGGAACCTT
TTCCTGTTGATTTGCATACTTGGAGGATCTGGATCAGACAGTGTGCTTATCGTGGTCAGAAAATGCCTTTATTTTATTC
AAGCAGAGATCTGGAAGGTAACAGAATTTCCACACATGCTTTCCATCTCTGTTTCAAATCTTTGGCAGAAAAA
ATCTGTTATTTTTAATTAGATAAAATGCTATGTTAAATAATTGAAGATTTAATTTCTTACTCTGTCAATAGAAGGCAGA
GTAGTTTAAAGACTTGAGAAAAGAAAAACAATTGTTTRCTTCCTATTAGAAATCACAAAATCCTCTCTGAAGTAAACA
TACAAATCTTTGTTTTTCTGCTTTCAAACAGGGAAAATTTTTTCAGTGCCTAACAAACAGGTGTTTCATGATTAGATTA
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GGAATTCCTGTTATGACTAAGAGTTGATGTAGTTAATTTAGTATGTCTAAAGTTGGCTGTAATCTATCACCACAGAAGA
GCTGGAAGAAATGATCTTAAATCACAAAGGATTTAGTAAAGACCTACGTGAGCTGTGAAAGATGATTAACTATGAAAT
AGTTCTCAAAGGAAGTTATAAATTCCTTGCCACTTGTGAAATCTAAAACTACATAGACAGTTATGTTTCRTATTAGAA
AATAAAGATTTATCCTGCCTTGGCACAATGTATTGAATTGGCATGTTGTTCTCAGTAACAAAGTCCCAATTTGCTTCAT
CAGCTTTAATTCCCTATTTACTCCATACATACTCTATACTCTATGTGTTAGTCTGAGTCTTGCAAGAAGCCCATGCCAA
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CCAAAGGCATATGGGAGTCCTCAAGCCAAAATCAGCCATCAGAGGAATCCTGTTTCCCAGGAATGTGTCTGCCACAACA
TGCCCTCTGTGCTCAGTAAATACCAGGAAGCAGGGCATGGGAGGTATGGCCTTAGCTAAAATGTTGCAGTGAATTTTCAG
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GCCCCAATCAGTTACCACTTCCTTCATGAAGGTTACTCTGTACAACCTAGTGGCTAATAATTACCATTCTCTTTTGTAA
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CAAAATTGGCCTAAAGCCAGATTATATGGGGCATCGGGTCTTGGCAAGTAGTTTGGACTTTATTCTATGAGCCCAATCC
AATTCAGAACATCTCAGAGTCATGTGGGCCAGGCTGAATGACAAAAATATTGATGTATATAATTCTGTTATGGCATACT
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GAGTGTGCTTTATAATATTTGGAAAATAGTTATAACTAGTTGAAGCATTCACCAGAATGTCACTTGACAATTCTAACTT
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CTTATTGATACACATTAAGAGCAAGTTGCAGACATGATACTCCTTTGCTTCTAAATACTTAAATAATTCCTAAAAACAT
GGAATTATCATACATAGGTACAGTTCAAGTTGTTAAATTAGCAAATTAACATGGATAAAATGTTGTTATCTAATCTACA
GACTTTATTCTAAATTTCAATAATTGTCCCAATAATGTTCTGTGTAGCCAATGAAATTACAAAATCATGCTTTTCAATC
AGTTGTCTATGCTCTTTAATATCCTTTAAACTGGAGCAGTTTCTGAGTCTTCGTGAAAGACTATTTTCATGACATTTATA
TTTTTGGAGAATTACATGCCAATTGTTTTGTACACTCTCTTGATTTATTTGGGCTGCTATAACAAACTACCATAGTCTT
CATGGTTTATAAACAACAGAAATTTATTTCTCATGGTTCTGGAGGCTGAGAAATCCCATGATCTGTGCTAAGCAGATTT
TCTGTCTGGTGAGGATCAGCTTTCTAGCTCATAAATAGCTGTCTCCTTGCTGTGTTTTTACATGGCAGAAAGGAAGAGA
GAGTTCTCTGGGGTCTCTTTTATAAGGGCATTAATCCCATTCATGAGGGCTCTGTCCCCATAATCTCATCACTTCCAAA
AGGCCACACCTGTAAATACCATCATATTGCTGATTAAAGTTTCAACATATGAATTTGGAGGCAACGTAAACATTCAGTCT
AAAGCATAGACTATCTTTCAATTTGAGTTTATTTTGTGTTTTCTGGGGATGATAGTCAGGTCATACAACTTGGGCAGGA
ATATCACAGTAGGGATGCTAAGTCCTTATATCCAGAGGTAGATGCTGTTGTTTAGTACCATTACTGACAATTTTAGCTT
TGATCATTTAGTTCAAGTGGTGCTACCAGATTTCTCCACTGTAAAGTTACTATTTTCTCCTTTGTAATTAAGGAAGCA
TTTTGTACCATGATAATATGAGAGCATGTAAATATGCTGTTATTCCTCAATTTCTTACCCAATAGCTTTGGAATCCGTT
GGTGATTCCTGTCTAAATCAGTTACTTGTGTTATTATGATGGTTGCTGTTTTTCATTGGTTTAAATGCTGCCGTAACAGAT
ACCACAACTGGGTAAATTTATAATGAATAGAATTTATTTGGCTCATGGTTCTGGAGGCTGAGAAGTTCAGGATCAAGGG
GCTGCATCTGGTGCAGTCCTTTTTCTGCTGCATCATGACATGATGAAAGGTATCACATGGGCAAGAAATAGGGAGAAG
GGGGCCAACTCATTTTTTTTTTTTTTTTTTTTGGAGACAGAGTCTCGCTCTGTCAACCAGGCTGGAGTGCAGTGACTC
CGCCTCGGCTCACTGCAACCTCCACCTCCGGGTTCCGCATTCTCCTGCCTCAGTTTCCCGAGTAGCTGGGACTACAGG
CACCCACCACACGCCCCGGCTAATTTTTTGTATTTTGTTTAGTAGAGACGGGGTTTCAACGTGTTAGCCAGGACGGTCT
CCATCTGCTGACCTCGTGAGCCACCCACCTTAGCCTCCCAAAGTGCTGGGATTACAGGCCTTAGCCACCGCACCTGGCC
CAAACCTCATTCCTTTATAAGGAACCTACTCCTATGATAACAGATAAAGTAATCCATTCATGAAGGGAATGCTCTCATGA
TCTAATCATCTCTTAAATGTCCCAACTCTTAACACTGTTGCGTTTTGGTCTAAGCTTCCAAGATGTGAACCTTTGGGAGA
CATATTCAAACCATAGTGGTTTGCTAAGGGGAAATCTTGTAATTCATCATTTCTTCTCTTTTATTAGTTGGCTTTCT
ATTATGCATAAAAATGTTTCTTCGTGCTTGTCTGTGTGGAGTCATGAATTCCTTATTTAGCAGGTTGTTATCCTTTA
ATATTGTTGGTTAATATCATTATTTATCTTAATCCTCATACTGTCTGGATTGGACAGTACAAGGTAAGTCTTAGTTGA

Fig. 9.197

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CTCCCTTGTCATTTTAAACACGTTACTATCATTCTTTGAGTACTTTTTTACTTTTTTGGTTTTAGATATGAGACTCTTGTT
CTTTTATCCAAGCCTTGGAATCAGCTACTTTTATAGGGCTCTCTGTTTCCTTTCAGTAAAGAATGGTTTTTAGAAATCA
AAATTTGGAGAGTAGGTGTGCTTTTGCTTCTAGGTCCTTTTAGTAGACATATCTAGGGGGAAGTGTATGTATGTACAT
ACTCATATATAAATGTGTGCATATATATGCATACCCACACATACCTATATCCATGGGTATATTCAGATATTTGTATACA
TTAAACCTAGGATTTTCACAATGATAATCCATCATTGCAGTGCTCATTCTGTTCTTCTCCATTTTCGTATCTGTAACCTCT
GTGAGAAGTTACGGTGAGAAGCCTGGCTCCCATTTATTCACAATATATTATATATTGCATAGGATTTTCTTACCTGTGGA
ATGTGGTTTTCAAAAATTACTTTCAGAATTACCCACTCACGCCTCTATGGAAAAAGAGAAGTAAGGCAAAGAAAAGTATT
TCAGTAAGTAGACTTTTAGCATTATTTATGTTTTATATGCCTAATTCTGTACTCAAAATTCACCTGGTTAGTCTTTTC
TCACAGCCACTTCCACTCACTTCACGGTTCGTACTTATTTTAAATATGGATCAGTTAATTTGTTTTAATATATAAAT
TTGTTTTTCTCTAGAGTTGAATCGATTAATCTGGAATGATATTTTCTGTTTCTATGTTTAAAATATCAACTAATAATTT
CCTTTCCTTAATAAAGAAATCTCTAGCACCATTAAAGACTCAGTTAACTGAGGTGTAGGTATCCATGTTTTCCCTAGA
GTATAGAGCTGTGATCTAAATCAGAAAATCTGAACAAGTTTGTCTTGGACCAAACCTTCAGGATAATCTGGAAACAACGG
GACAAATGTTAGTGCATGAGTATTTTCAAACATAGTTCTGAGTAAATTCCAAATGCATATGTGGGTTTTGTGGCTCTTCA
TATCTGGTTATTTGTACATGATCACATCTTTATTGTAGATAATAAAAGTTGGGTATAAAAGCAATGCCATTAATTATGT
ATTTTTTATGCTTTTTTGTTCAAAAAGGAATTAAGCAATCCTAACTTTAAAAATGGATAAAATGCCTATGTAAAGCTA
AGTTAGAAACACAAATGTAAAAGTGTATTTCTGACTTGGAGTTAAGTTTAGGTCTTAAAATATTAATAGCTTCCTCAA
GATACAGCAAATAGAGAAACCTGATAATATTTATGTCTTGAGAGAAAATTATCTTTCTATAATTAATTATAAGACTTTT
CATTAGGGTCTTTGTGTAAAGTAAAGCAATAATATGTTTAAATAGTATTTTTATGTGAGTTACATACATGTACATTATTA
ATCATGAATAAATAGTGATGTTCTTATACTTAGTCTTGTATAAGTCAAAATATTAATCATTATTCCATGAAAGTCTT
CATATAGTAATTTAAATCATTGTTAAGTCTGGTTACTTGATGTTGAGAAAGAGCTTAGTACACATAGTAGAATGACTTG
CCAGCCTTGCCATTAGGTTATCTTCTCATACTACTTACCTTAAGTTGGCAGGAGCTTGATTACAGAAGATAACTGT
CATTTATCTCTTTGTGTGGAGTGCATATATTTGTGGGATGTCTATAGGAGTATTACCCAAAGGGTTTTGTATTACAGAATT
ATAGTTAACAAAATCACTAGGAAATGTTTATGGAGGGAAGATAACCACAGATATATCTGATTACCTATTTGGAGGACTA
GCAGTGGAATTTCTTGCTGTTTCTCATGTGGCCAAAACCTAATAAAATAAAATAAATTTAGAAGAGTTTTGTGTACCAA
GATAATCCATATATTTTCAAGTTGGTATGGAAAAACATCTTTTCTTTCAGCCATTGTTACTATGACACTTAGACACTCTAG
TCTACAAAATTATCCTAGGTTTAGCAATATTTGGAATTAATTTTATTGTTCTTCTGATTGCAGAGTTATAAAAATAA
GGATTGAGTTATTATGAGAGGCCGATTGTGTAATACATATCATGAGTTAAAAGTAAACGTTGACTTTTATTATAAATTGC
CTTTTGCCACATGAGAGAAAATAAACTAAGTGAATAAGACTGTCTCTAAGTCTTCTCTGAGATTAATTTGACAATTTAC
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TGGTGTTTAAACAGGCCAGGTGCGGTGGCTCACACCTGTAATCCCAGCACTTTGGGAGGCCAAGGCGGATCACTTGAG
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GCCGAAATGGTGCCACTGCACTCCAGCCTGGGTGACAGAGCCAGACTACATCTCAAAAACAAACAAAAAACTAAAACAG
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TAATTCCTTACACATTTTAAAGTATATGGGCTAAAGTATGAAACCAACTCTGAGTAATAGGATTATAAATATTTCTTTAT
CTCCTTGATTTTATATTTATTTATACCTTAATTAAATTTTGAAATAATGTTGTTACTTTTACAATAAGAAGTTGTGAAA
AACTATTACCATGAAAATAATCTAGCAAGAATTTGATTTAACATTGCTCATTTCTTGAAAGCAAGCCAAGACATTTTTT
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TTGTGCAGGCACCTCAGTTTTAAGTAATCATGGGATTATTTATTTGTTGATTTAGCATTGTTCAGTACTTATGTGCTT
AATAGCACTTGAACATACAAAGAGAAATGAGATTTTTTGGTCCCCAAGAATTAATAAATTTAATGGAGGAGATAATCATAA
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GGTCACACCTAGGAGGGGGCCAAAGGGGACAACTCACTGGGTATTGGAATGATCTGGATCCCTGGATTACTTGGTTGGTG
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AAGTAGGGTGCTTTTTTGGTGGATCAAGAGAGTCCAGATCACTAATATATGGATGGCGTATAAATTAATCACTGTGTAA
GACTGCAAAATGTCAGAGAAACATTTTAAATATGTTCCCCCAGCATTTGCCTTTCTGTACATATGTCTTAGAATTATAA
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GATTACTAGAGGCTGGAAACAGTAGGGAGAAGGGGAGGATAGAGAAATATTTGTTAAGAATACGAAATGACAGCTAGAT
AGGAGGAATTAAGTGGTAGTGTCTATAGCACTATAGTATGACTATAGTTAACAACGATATATTATATATTTTCAAATA
CTTAGAGGATATTGAATAGTCCCAAAGAAATGAAAAATGTTGGAACATATGCTAATACCCTGATCTGATCACTATGCAT
TGTATGTATTGAAACATCCCTACGTACAATTATTATGTATCAATTTAAAAAATATCTAAAAATATATGTGTACCTAAGT
GTATACACCTTATATATCCTTTGATTTCTTATGAAACACAGCATAGTTTGAGATTTAAATGAATAAAATGGTTCTTT
ATGCTCTGTAAATTAGAAAATCCTTCAGCTTTTAGAAGCTCTCTGTTGGTTTATTAGTTGATCACAAATTAATATGTC
CCTTTGGTTTAAATCTGTATCATAAAACAAAAGTTGACTGTTCCCAAACATGTGCTTGGATAAGAGAAAGCTGTTAATAA

Fig. 9.198

[illegible]

Fig. 9.199

Fig. 9.200

AGAGAGTCAGGAGGGTTGTAGGGGCAAATTTCTAACGGGTTGTGGAATCTATCATTTGCTAGGACCTTTTACAATTTCAGTT
GCTTACTCTTTTTTAAAATCTGCCATTTCAGCAGTCTTTTTTTCAGTACCAGTAAGCCACGCTATGCATTCTATGAATAAAG
TGCACAGCAACAGCATCCAAATGCAGTGACGGAACCTGTATGGCTTCTCCTTGGAAATAGCTTTTAAAAACCTGCTTAGGC
AGGGACACTTTTATTGCTTGCTGAAAAATTCCAAAGATTTGTGACTATGTTTAAATAAATTATCAAGAACTATATAGAAT
TCCTTAGACCTTTTTTCTAAAATAAACTTTTGGAGATTTTTTGTGTTGTTGTTGTTACTATTATTTTTTAGAGAAAGGAC
AATTAAATTATCTTCAGATCCCAGGGAATGGTAGAATTAGTCTTTTTTGGCACTGATAGTAAGGTAGGTCATGTCTTAGT
CCATTTGGACTGTTACAATAAAATATCATAAACTCTGTAGCATGGCTGGGCACAGTGGTTCACGGCTGTAATCTCAGCA
CTTTGGAAGGCCGAGGCCGGTGGATCACCTGAGGTCAGGAGTTCGAGACCAGCCTGGGCAACATGGCGAAACCCCATCT
CTACTAAAAATACAATAATTAGCCGGGCATGGTGGCGCATGCCTGTAGTCCCAGCTACTCGGGAGTCTGAGGCAGGAGA
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CTCTGTCTCAAAAACAAACAAAAAACTGGGTAGCTTATAAACAAATAGAAATTTATTTCTCATAGTGCTGGAGGCTGGGG
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GTCCTCACTTGGAAGAAGGGGCAAGGGAGCTATAACTTGGAAGAAGGGGCCTCTTTAATATGTGCATTAATCCTGTTCA
CAAGGCTGGAGGCCAATGACCTAATTACCTCCCAAAGGCCCTACCTCCTAACACTATTGCATTGGTGATTAGATTTCA
ACATTTGAATTTTTGGGAGGGTGCCAATGTTTCAGATCATAGCTGGTCATTTTCAGCTTATTTGTACTCTCTAAGTTATGCT
GCAATAATTGTTGGCTTCTGTACATGGGCAATATGGAAATTAAAGCAATCATAACATTTTGGTAGAGGACCTGAGGAGA
GATTATTCTGAATGTGATGAAGTTATAGTCTTTTTTCTCAGGTGGATTAGATCACTTTTTTGGATAGAAATTTTATGTTG
GGAAAAATAAAAAATGGTTCAGTGACCTGAGGCAGAAAAAGTCTGTTTACAGGAATATAAGATAATTACTCCATCAACTC
ACAGTTGTTTGGAAATGTGAGGATCAATAATAGTCCCTCAGAACATTGTAACCTTACTCCAGCAACTATGCTAACATAAAG
CAGAGAAGTCTGAAGTCTGTCCATTAGTATTGGTGACAAGGTCATGGTTGCTTTTTAACTTTGCTTGCCAAAATACCTG
GAGGGAACTCAATGTGCATACTCAAGTTTCTGCTGTGGCAAATTGACTGTTGCTACCCCAAGGAGACCCTAGGAAGACAA
TCAGTTCATGTCTGACTGGAGACATTGCAACTTTATTTGTATGAACATTTTGGGTCAAATTTTCAGGGGGACATGTCCT
ACAGAGGGGCAGAGGGTCATGATAATGAAAAGAATCATTACGTTATATTGGTAATTAGATGTATGACTCTACAACTTG
TTTTTCTTGGGTGTTAGTAACATTCTTTTTTGGAGATAGATGGCATTTTACAAAGTCTATCCACAGAAAACCTGGAAGTA
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ATTTCTTTTAAAGTATAGAAAGGTTAGACTTATCAACCTTATTTGCAGATACTATTTATGTTTTTTCATGTTTTCTCTCTG
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ACCTCTCTTTATGAATGAAATTTTTTCACAAAACCTCTGATATATAAAACAGGCAAAATGAAACCATTATGGCAGGAATGA
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GAATCAACTGTTGACTACTTCTGAAACCTAAAATGAAATAATATATATGAAAGCTGTTATTTCAGTGATTTATACTTATC
AAAATATACTTGTCAAATATTAACCTGATTTGTGTTTGTCAAATAAATATATTCATACATAGCACCTTGGTAAATCAC
AGATCTAGTTGTATTGTAGGTGCCTAGACCAACATGACAACTTGATTATATGGCTTGAGATATTTACATCACTCTTACC
ACAAAAACGTCAATAAACAGAGCTTATTAATCTTTTGGAGGAGGAGGAAATGAAGGTGTAGGAAAGCATGTATGAAGTTC
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CACTAACTCTATAAATTTCTTGTCATATTAGGCCTGTTTCTAAAGCCAATCAGAATGAAGGAAGAGCTTTTAAAAAAT
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CATTAGAAGAAAGTAGAGGAAATGGCTCAACCTTGCCAGAGTACAGCTGGTTGGAACCTTGACAAACGGAACGTGGCC
CAGGGTAAAGAGCAAGAGGCTGGGGGAAGCACGTAATGCTTCTTAAAGCCTTTGATCTAAAATGACAGACTGTCATCTC
CACTCACATTCATTAGCCCAAGCAAGTTTTGTGGCTAACCAGACAATGGGTCTCAGCAATATATCCCCCTCCCAGAA

Fig. 9.201

CGCCTGAACATTCAAGTGGCAAAAGAGTGAATTGTTGAGAACAATAATATAGTCTATAATAGCATATAATGTTTCATAAA
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TGACAAGTTACACTTTCTGAGACAATCTTCAATTTTAGCAATTAAAATAGCAAAGAATTATTTATAACAGTATTCA
TAACATAGCAAATGGTCATATTCCTTATAGAGACAGCACTTCCGAGTAGTCTTTATTTTGTTTTTTCAAATTTGTCCTTT

Fig. 9.202

Fig. 9.203

TTAATTTTAATAGTAAACTCTTTTAAGGAATGAATGGCCACAGGCAGAGACTAATGGTGTACTTTCAGTGTCTCAGTG
AGTGCTTCACAGATTAGGTTTCCAAGAGAAGAAGATTGCCTCTGAGGAAGAAGAGATGGGGTAGCAAGAGGCATATATG
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TATT
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GGTAGAAGATACGGTTAAGATGAATTTTAGTTATTTAAAGTGTTGTCATAACCTGAACATTTTACAGATATATTTAAAC

Fig. 9.204

AGGTATAGGTATATTTAAAATCACTATTTTAAATTAATAAAAAAAAAAACTTTGAAAAATCCCTAATTTCAAAAATATATT
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AAATTCTCTAGACTCTATCTCATGCATTATTACATCTCTCTTGGCATACTTCTCTTCTTGAAGTGCATAACATTTATAT
ACCTTCTACAAATGTTGGTATTATTGATATTGAGGGAGGGGGCAGACCCAGGTCTATGGCCTTATGTATTATTAGTGAA
TTACAATTTCTTAGAGTAATAGGCCAATGATTAAAGTCACATGTTAGGAAAACCTTAATCCACTAGGGAAAACCTTCCAA
TATTTTTCAAAGTAATTTACTACATTTGGATGCAAAATTATATAGGTGTTTTTGGTGGTATCCATACTTGCTTTCTTGTC
AACCTGAAGAGATGTAAATGTAAATCGATGCTAATTGTACCGTGCTTACACAAAGAGCTAATTTGAGTAATATCCTATC
CCATCAGTTCAATTTTACCATAACATTTTTTTCTTTACAAAATTGTTTTTAAACTCCACTGGAAAAAATATATAATAA
ACTACATGTAAATTACACAATTTAGTAAATTTTGATATAGGTATACACATCTGTGAGAAAATTACCAAACTGAGATAG
TACACAAATATATATCACCCCCAAAAGTTTCCTCGTGTCTTTGTCAATTCCTTCCCTCAGCTTTTTTCCCAGAAATTTAC
CTTCTCAGGCTGATCAACACTCAGCTGAAGACTCAAGAGGGGGCTGTGCAGATGTCCCTGAGTTCCCTCTCTGGGCC
ACTGTCTCCTCTCTTTTTCAGCAAATTTCTAGTTACCTTGACCTCCCTGAATTTCCAGCTCCATCTCCTCAACTTAGGGAG
ACTTTTCAGGTGAGGAGTGATCCCTCCTCCTTATGCTGGGACCTGGAAGCTCTCCAGCATGTACACTGGGGCAACCATA
GGCGTCATCTCATTGTATTTCATTTTCTCTAATCACCATGCTAAGCTAACTGTTGCCTAAGGTCTGAAAACGTGTTGTTT
ATTCATTTTGTCCAGTTTTTCCATTCCTTTCATGAGCAAGCATAAACCTGACTCCTTTGGGCAAGAAGCAGAAGTCCCTA
TAGTGACTTTATTTACTTTTATTGTTATTGTGTCAGCATTTGGAAAAAATGTAGAGATTGACTATTTTATACTATTTAA
GTAATTTTTTGTATATTTGATTTTACCTAAACAAGAGTTTATTAGAAGTATTTTGTATACAGCTTATAACAAATAATT
GTACAAAATTAGAAAAATCAGATGAACAAATTTTTTATAAGACTTCTCGTGGTTATAGAATGACTCTAACACCTTGTA
GTGTATTTTTTAGGATTTTGTCTTAGTGTCATGTGTTAGTCTGTTTTCATGCTGCTGATAAAGACATACCTGAGACTGG

Fig. 9.205

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GAAGAAAAAGAGGTTTAATTGGATTTACAGTTCACATGGCCGGGGAGGCCTCAGAATCATGGTGGGAGGTGAAAGACA
CTTCTTACATGGTGGCGGCAAGAAAAAATGAGGAAGATGCAAAAGTGGAAACCCCTGATAAAACCATCAGATCTTGTGA
GACTTATTCACCACCACGAGAATAGTATAGGGGAAACTTCCCCCATGATTCAAATTATCTCCACCGGGTGTCCCCAC
AACACATGGAAATTATGGGAGTATAATTCAAGATGAGATTTGGGTGGGGACACAGAGTCAAACCGTATCAGTGTATATG
TTTAGAGGGACATAGAAAACATATATAAGTTTAATAAGAATATATACATATTGTTATTAAATGAAAACATTTGAAATTAT
AAAAGTAGTAAATATTTTTTAGTAAAATTAAATATATATAGTAAAAAGTGAAAGCTGTTTTACTTCCATTCCCAGAG
AAAACTTTACTTAAATTTTATATAGTTTTTTCCAGAGTTCCTTCTACACCTGAAAAACACAGTGGATTATTTTTGTGTA
CAAAAGAAAAGAAATCATTCTATATGTATCTCCTTTACTTCCACTCTTTTTCTTTCCACTTAATTCATACAGATAAACT
TTACCATTAAATGGCCTCTGGTGATTCTTTTATAAGAATGAACCATTAGATAGTTAACCAGTTTACTAATGACAGACA
TTTATATTGCTCCCAAGTTGTTGCCATTGTTCAATTTATTTATTTCTTTAGTATTAATTCCTAGATAAGTTGCTACATCA
AAAATAATGTACATTTAATATTGCAATAGTTACAAGGAAATCTGTACTTTTTTTCAGTAATATATTAGTGTCTACTTTCA
AAAAGCTTTAAATCTGAGCATCATATTGTCTTCTAAATCTTTGACAAAAATTAATGTAAATGAATATGCTATTTTAAT
TGTCATTTTATTTATTTGATTATTAGTAAGATTTAGATTTAGATTTTCAGTCTTTTTTATTAGCCATTTTCAGAGGGATGAGC
ATTCTCTGCCTATGGACTTTGCCTGGTTTTGTATTGATGTGTTTCATTGTTTTCTTATTTTCTCATTTGTAAACATCATTT
TAAAGTGCTTTTTATTTCTGGACAATGTTTTTACATTTTTTAAAGACTCCTATTGTGATTTTTTATTAAACAAAGTTAGAT
YTATAGAAAAGCTGGAAAGAAATTTGAAGTTTTAAGCATATTCGCTCTCCCTTTCCAGAATATGTGGCAAAGGAGGGAAA
AGAAGGGATTCTGAAGGAACACACGGGAAGTCGGCAGTGTGGGAGTGATGGATATGTTTCATTCCCTTGCTTGTGGTGAT
GGTCTCACTGGTGTATACATTTGCAGTAATTGTCAAGTTGTACACTTTAAATATGTGCAGTTTATTTTTATCTCAGTTA
CATCTCAATGAACTTTTTTTTTTAAAGTGGCACCAACATACATTAAATAGCTTTTCAGCCAAAAGTAAAAACTGAGTTTG
GTCAGAATATTTTTGTGCTATTTTTGGTTAGATTTCTTCTGTCACAGTTATAATGGCTTTCTAAATAAAAYTAGGAAGCC
GGAGAAAGTCACTTTAACCTTCAAGAGTAGGGAACTGAACCCATAAAAAAAGTGCTTCCAGGTCAGACTCAGCCTAAA
AAACCTGAAAAATAAATGTGACTTCTGAGTAGCAGTATATGGGAAAAATTTTCTATAAATCTCTGAGAATGAAGTTGAC
TGGAGACTACACAAGAGACAAAGAGATATAGAATTGCCTTAAATCTGAACCATGGAGTTACAGAATGATAATGGCATT
TGATAAATTTTTAGAGAACTCTTGATAAACAGGCACACTCAAAAATAGCTTTTTGTGTTGTTGTTGTTGTTGTTGTT
TTCTGTGCAAGAGGTGACAATTGAATGAGGTTCCAAAAGTAGTGTGGGCTTGCAATTGCACACATATTATCAGGCAGGA
TTTGTGCAGTAGAGAAAGGTGACTGAAGAATGTAGTGGAAGGACATGCTACTGGAAGAAGCTGCCAAGTGACATACACA
GGGCTGTGAGTCAGCCAGGGTTGAGGCTCTGTCCAGTGGGACAGCAGGCTCTGTCTCCACAGTGGGGACTGCCTGAGAG
CTGGGTGCTCTGGTTCAGCCATTGTAGCACAAAGGCACCTTGATAAATTTCCGTGAAATGGCTTGAAACCTACTTTGCC
ATAAGAATGAGATAAATTTTTAATTCAAAACCTTAAATATCTTGATTAGAATAAGGAGGAGGAAGAGAAAGAACAGGCA
ATAAAAAGTTATTTATTTGATTGATTGACTTAATGCAGCTAGCTGAAGGGTGAGAGGAAGAACTCCGGCTCCAGAGTCTG
AAAGCCAGGGCTCAAGTTTCAATTTGGGCATTTCCAGCTTTGAACAACAGAAAAACACTGTTTACCTTTTCAGAACCT
CAGTTTCCCTTAGATCTGTAAATTAGCAATAAAAACTAATGTGCCTTCCAAGGTTATGGTAAAAATCAAATATCTTATGC
CTGTGTAAATCTTTTTCAAAAAACAATAGACACTGCAAATATTGGGCATTCTTATGATGATGTTTATTCTTCACTGGGA
GCATTGATGGATTGATTGTTACTTTTTCAATAACTTTTTCCATATTTGCTCTAGTTTTAAATTTGCAAAATTTTAATTCAG
TATTGTTTATAATAAGACAAAAGCTCTTCTTTAAGGTTGGGGCATTAATGTTAAAAAAGAACTGTTACATCA
AATGTACATCAAATGCAGTGAAGTGGACAACAACCAAGAACTTCTGTGTAATCCCATCCCATCACACACACACACA
CACACACACACACACACACACACACTATAGACACTATATTTAAATTTGGCAGGGGTGATATTATGAACTCCTG
TAAACCAAATTTGATTTGTGTTTGTGAGGAATGTGTGGGTCTGTGTCTGGAATGCTATATATTTCCACTGTGCACTGC
CCATTGACCATTTCTACTAGGATGCCTTGAAGGCACCTCAGACTCAACAAGTTCAATTTGAAATTCATTATGTACTCCC
CAGTTGTTACGCTAGGAGCCTTCAAGGCCACCCGTGACTTTTCTCTCTCCTTGAGTCTCTCCTTAGCTTCCCATGGATC
AGTCTCTGTACCATGTTGATTCTACCTTCTGAGTATCCTTTCTGCTCTCCCTCATCTGCACTTCTGTTACCACTGCCTG
AGTTTTGGTTATGGGCTTCTTAGTCTTTTAGTCTTTTCTCTCCCCAACCAGCCATCGCCACCAGCAAGAACTATCTT
TCTAAAGCACCAATTATCTTTCCAAAACAGAAGATTTTCTCTTCTTTGTAGGAATTTAACTTTAGGGCATAGTCAGATA
AATTAGCCAAAATATATGTGAAGAAGCTGTTTCATCACAATATTGTTTATTATAGTAAAAAAGGAAAAAACCTTAATA
TCTCAAAAATGGATCAGTAATATCGAGACTATACACAATAGTCATTACATTGGTATTATAGATGTATAATAATTTGGAA
ATGATATTATAAAAAATTATTAAATAATGTGCATAGAAAAATTTGTGGAAATGTTTTATGCTATAAAAAATTTCTAG
GTAATGGAATTTAGTAAGAATTTTTCTCTTCTACTTCCCACCAAACAAAAGCAAAATAAACTAAATAGAACAAAA
TTCTGCTATGTAGCCCTTTACAAAAGATAGTGCTTCTGGCAAGAACTGTGACTGACCAAAATCCTGCAATTCAGTACC
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TCATTTTTTTCAAGAAAATTTGTTGTAGAATCTTTTATATAAAACACTATACACTGTCTTAAATCCACTATGTCTTATTC
CCTCTATTTACAAATGTGCTTTGTCTTACATTTCTTTAAAAAACAACAACAAAACCTGTGTTACACAGTATCTTTG
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AAGCTCCGTATCTCTCATTTAAGGTCCCTGATAGGAGTTACTTGTCTCAAACATGTGTGTTGGCTCTTTACTGTCT
TTTAAGTAGCATTTATGATGAATTCCTTTTTTTTAAATGGTGGTATTACAGTATCTGATGTTAGGCATAGAACTGCTCAG
GCAGCAAGTCACCTTTTCTTTTTTCCCTTTTTTGTAGACTTGAGAGACCAAGGCAGCACACACACACTCCTCCAATA
GCTTTGCAATTTGCATACATTTCTGATTCTGGTGGKCTATTCATCATTTTTTATAGAAATGCTTTTTATGAAATTCGATCTC
ATGGAGTGAGTTAGGAGTTTCTTTGTTTTTGTCTTTATGATAACCCATTGCCTCAGTAAATATCTTAATTATTCAGAG
AAAAAATAAACCCAACTGATTAGACCTAATCCCTTCTCAAATCACTTTACTTTCTTTGTCTATTTCTATCCATCCTCTC
TCTACACTCAGATGTCTTTGTTTTTATTATGAAATATATTTGGCATGAGTAACAACATTAAGTAATATGAATCATTTTAT
TTGGTTAAATGTCTACACTCAAAGATTAGTGGAGAATTTGGAAAATATAACGAAATAATACTGGAACTTTGTATTTTC
CCTAGCTACTTGCCATATTCAAAGATGTGTCTTTGAAGTGAAGAGCAGAATAGTACCTTATTATAGTCCATATTAAAT

Fig. 9.206

CATATTTATGGCTTTCAACAGAAGTATTGAAAACCAGATTTCAGTTGCTGCACCCATCAACTCATCATCTACATTAGGTA
TTTCTCCGAATGTTATCCCTTCCCTTGMCCCCAACCCCCGACAGGCCCCAGTGTGTGATGTTCCCTCCCTTTGCCCA
TATGTTCTCATTGTTCAACTCCCCTTATGAGTGAGAACATGCGGTGTTTGGTTCGGTTCCTGTGTTAGTTTGCTAA
GAATGATGGTTTCAAGCTTCATCCAAGTCCCTGCAAAGGACATGAACCTATTTTTTTTTTTATGTCTGCAAAGTATT
CCATGGTGTATATGTGCCACATTTTCTTCATCCAGTCTATCATTTGATGGGTATTTTTGTTGGTTCCAAGTCTTTGCTAT
TGTGAATAGTCCTGCAATAAACATACACGTGCATGTGTCTCTATAGTAGAATGATTTATAATCCTTTGGGATTATATAC
CCAGTAATGGGATTGCTGGGTCAAATGGTATTTCTACTTCTAGATCCTTGAGAAATCGCCACACTGTCTTCCACGATGG
TTGAACCAATTTACACTCCCACCAACAGTGTAAAAGCATTCCATTTCTCCACATCCTTTCCAGCATCTATTTTTCT
GACTTTTTAATGATCGTCATTCTAACTGGTGTGAGATGGTATCTCATTTGTAGTTTGTGATTGTCATTTCTCTAATGACCA
GTGATGATGAGCTTTTTTTCATATGCTTGTTCATATGCATAAATGTCTTCTTTTGAGAATTGTCTGTCCATATCCTTCGT
CTGCTTTTTGATGGGGTTGTTTGTCTTGTAAATTTGTTTAAAGTTCCTTGTAGATTCTGGATATTAGACCTTTGTCAG
ATGGAGACTGCAAACTTTTCTCTCATTTCTCTAGCCTGTTCACTCTGATGATAGTTTCTTTTGCTGTGCAGAAGCTCTT
TAGTTTAGTTAGATCCCATTGTCAATTTTGGCTTTCGTTGCCATTGCTTTTGGTGTTTTAGTCATGAACCTTTTGCCC
ATGCTTATGTCCTGAATGGTATTGCCTAGGTTTTCTTCTAGGGTTTTTATGGTTTTAGGTCTTAAGTTTAAAGTCTTTAA
TCCATCTTGAGTTAATTTTTGTATAAGGTGTAAGGAAGGGGTCCAGTTTCAGTTTTCTGCATATGGCTAGCCAGTTTTTC
CCAACAGCATTTATTAATAGGGAATCCATTCCCCATTGCTTGTTTTTGTGAGGTTTGTCAAAGATCAGATGGTTGTAG
ATGTGTGGTGTGTTTCTGAGGCCTCTGTTCTGTTCCATTGGTCTATATACCTGTTTTGGTACCAGTAGCATGCTGTTG
TGGTTACTGTAGCCTTATAGTATAGTTTGAAGTCAGGTAGCTTGATGCTTCCAGCTTTGTTTTTTGTTTGTGTTTGT
TTTTCTTTTGCTTAGGATTGTCTTGGCAATACGGGCTCTCTTTTGGTTCATATGAAATTTAAAGTAGTTTTTCTAA
TTCTGTGAAGAAAGTCAATGGTAGCTTGATGGGAATAGCATTGAATCTATAAATTACTTTGGACAGTGTGGCCATTTTC
ACTATATTGATTCTTCTATCCATGAGCATGGAATGTTTTTCCATTGTGTTGTCTTCTCTTATTTCTTGGGCAGTG
GTTTGTAGTTCAACTTGAAGAGGTCTTTCACATCCCTTGTAAGTTGTATTCGTAGGTATTTTCTTCTYTTTGTGCAAT
TGTGAATGGGAGTTTGCTCATGATTTGGCTCTCTGTTTGTCTATTATTTGTGTATAGGAATGATTGTGATTTTGTACA
TTGAGTTTTTATCCTGAGACTTTGCTGAAGTTGCTTATCAGCTTAAGGAGTTTTTGGGCTGAGACGATGGGGTTTTCTA
AATATACAGTCATGTCATCCGCAAAGAGAGATAATTTAACATCCTCTCTTCTTCTTATTTGAATATATTTTATTTCTTCTC
TTGCCCTGATTGCCCTGGCCAGAACTTCCAATACTGTGTTGAACAGGAGTGGTGAAGAGGGCATCCTTGTCTTGTGCCA
GTTTTCAAAGGGAATACTTCCAGCTTTTGCCCATTCAGTATGTTGTTGGCTGTGGGTTTTGTCATAAATAGCTCTTATTA
TTTCAAGACATGTTACATCAATACCTATTGAGTGTTTTTAGCATGAAGGAGTGTTGAATTTTATCGAAGGCCTTTTCTG
CATCTATTGAGTTGATCATGTGGTTTTTGTGATTGGTTTCATTTATGTGATGGGTTATGTTTATTGATTTGCATATGTT
GAACTAGCCTTGATCCAGGGATGAAGCCGACTTGATCGTGGTGGATAAGCTTTTTAATGCGTTGCTGGATTTGGTTT
ACCAGTATTTTATTGAGGATTTTCACATCGATGTTTCATCAAGGATATTTGGCCTGAGATTTTCTTTTTTTGTTGTGTCT
CTGCCAGCTTTTGGTATCAGGATGATGCTGGCCTCATAAAACGAGTTAGGGAGGAGTCCCTCTTTTTCTATTGATTGGA
ATAGTTTCAGAAGGAACGGTACCAGCTCCTCTTTGTACCTCTGGTAGAATTTGGCTGTGAATTCATCTGGTCTGGGCT
TTTTTTGGTTGGTAAGTTATTAATTACTGCCCTCAATFTCAGAATTTGTTATTGGTCTATTTCAGGGATTCCACTTCTTTC
TGGTTTAGTCTTAGGAAGGTGTATGTGTCCAGGAATTTATCCATTTCTTCTAGATTTTCTAGTTTATTTGTGTAGAGGT
GTTTATAGTATTCTCTGGTGGTAGTTTGTATTTCTGTGGGATCAATGATGATATCCCCTTTATCATTTTTTATTTATKG
TGTCTATTTGATTCTTCTCTCTTTTTCTTCTTTCATTGATCTGGCTAGTGGTCTGTTTTGTTAATCGTTTCAAAAACCAG
CTCCTGGATTCAATTGATTTTTTCAAAGGGTTTTTCTGTCTCTGTCTCCTTCAATTCTGCTCTGATCTTAGTTATTTCT
TGCCTTCTGCTGGCTTTTGAATGTGTTTGGCCTTGCTTCTCTAGTTCTTTAATTGTGATGTTAGGGTGTCAATTTTAG
ATCTTTCCTGCTTCTCTTGTGTGCATTTAGTGCTATAAATTTCCCTCTAAACACTGCTTTAGCTGTGTCCAGAGATT
CTGGTACATTGTATGTTTGTCTTATTAGTTTCAAAGAACTTATTTATTTCTGCCTTYGTTTCGTTATTACCCAGTAG
TCATTCAGGAGCAGGTTTTTTGATTTTCCATGTAGTTGTGCTGTTTTGAGTTAGTTCTTAATCCTGAGTTCTAACTTGA
TTGCACTGTGGTCTGAGAGACTGTTTGTATGATTTCTGGTTTTTTGCATTTGCTGAGGAGTGTTTTATTTCCAATTAT
GTGGTCAATTTTAGAATAAGTGCAATGTGGTGTGAGAAGAATGFATATTCTGTTGATTTGTGGTGGAGAGTTCTGTAG
ATGTCTATTAGGTCCACTTGGTCCAGAGCTTAGTTCAAGTCTTGAATATTCTTGTAAATTTTCTGTCTCGTTCATCTGT
CTGATATTGACAGTGGGGTGTAAAGTCTCCACTAGTATTGTGTGGAAGTCTAAGTCTCTTTGTAGGTCTCTAGGAAC
TTGCATTATGAATCTGGGTGCTCTTGTATTTGGGTGCATATATATTTCAAGTAGTTAGCTCTTCTTGTGTCATTGAACCC
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CAACACCTTTTTTTTTTCTTCTGTTTGTGTTGGTAAATCTTCCCTCCATCCCTTTATTTTTTGAGCCTGTGTGTGTGTTG
CACATGAGATGAGTCTACTRAATACAGCACATTGATGGGTCTTGACTCTGTCTAGTTTGGCAGCCTGTGCCATTTAATT
GGGGCATTTAGCTCATTTACATTTAAGGTTAATATTGTTATGTATGAATTTGATCCTGTCAATTATGATGCTAGCTGTTT
ATTTTGCCCATTAGTTGATGTAGTTTCTTCATAGTGTGAGCGGTCTTACAATTTAATATGTTTTTGCAGTGGCTGGTA
TCAGTTTTTCTTTCCATATTTAGTGCTTCTCCAGGAGCTCTTGTAAAGCAGGCCTGGTGGGGACGAAATCCCTCAGG
ATTTGCTTGTGTTGTAAGGATTTTGTCTTCTTCAATTATGAAGCTTAGTTTGGCTGGATATGAAATTTCTGGGTGCA
AATCTTCTCTTTAAGAATGTTGAATATTGGCCCCCACTCTCTTCTGGCTTGTAGGGTTTCTGCAGAGAGATCAGCTGT
TAGTCTGATGGGCTTCCCTTTGTGGGTAAACCAACCTTTCTCTCTGGCTGCCCTTCACATGTTTTCTTCAATTTCAACC
TTGATGAATCTCACGATTATGTGTCTTGGGGTTGCTCTTCTCAAGGAGTATCTTTGTGAGGTCTCTGTATTTTCTGAA
TTTGAATGTTGGCCTGTCTTGTAGGTGGGGAAGTTCTCCTGGAAAATATCCTGAAAAGTGTTTTCCAACCTTGGTTTC
ATTTCTCCCATCTCTTTCAGGTACACTAATCACATGTAGGTTTGGCTTTTTACATAGTCCCATATTTCTTGGGAAGCTTT
TTTCATTCTTTTTCTTCTTTTTTTATCTAATCTTGTCTTCATGCTTTATTTCAATTAAGTTGATCTTCAATCTCTGATA
TCCTTTCTTCCACTTGATCAGTTTGGCTATTGATACTGTGGTAAGCTTCACGATGTTCTCGTGCTGTGTTTTTCAACT

Fig. 9.207

CCATCAGGTTATTTATGTTCTTCTCTAAACTGGTTATTCTAGTTAGCTATTCCACTAACCTTTTATCAATGTTCTTAGC
TTCCTTGCATTAGGTTAGAACATACTTTTCTAGCTTGGAGGAGTTTGTATTACCCACCTTCTGAAGCCTATTTCTGTG
AATTGATCTAACTCATTTTCTGTCCAGTTTGTGTTCCCTTGCTGGCGAGGAGTTGTGATCCTTCGGAGAAGAAGAGGCAT
TCTGTTTTTTTGGAAATTTCCATCCTTTTTCGACTGGTTTTTTCCTCATCTTCATGGATTTATCTACCTTTGTTCTTTGCTG
TTGGTGACCTTTAGATGGAGTTTTTGTGTGGTCATCTTTTTTGTGATGTTGATGCTATTGCTTTTTGTTTGTAGTTT
TCCTTCTAGCAGTCAGACCCCTCTTCTGCAGGTTTGCTGGAGGTCCACTCCACACCCCTGTTTGCCTGGGTATCACTAGC
AGAGCCTGCAGAACAGCAAAAATTGCTCCCTGCTCCTTCTCTGGAAGCTTCGTCCAAGAGGGGCACCCAACAGATGCC
AGCCAGAACTCTCCTGTATGAAGTGTCTGTCAACCCCTGCTGGGAGTATCTCCCYATCGGGAGGCACAGGGGTCAGGGG
CCAACCTTGAGGAGGGAGTCTGTTCCCTTAGCAGAGCTTAAGGGCTGTGCTGGGAAATCTGCTGTTCTCTTCAGAGCTGGC
AGGCAGGAACATTTAAGTCTGCTGAAGCTGTGCCCAAAGCCACCCCTCCCCCAGGTGCTCTGTCCCAGGGAGATGGGA
GTTTTATCTGTAAGCCCTGACTGGGGCTGCTACCTTTCTTTTCAGAGATAACCCTGCTCAGACAGGAAGAATCTAGAGAG
GCAGTCTGGCTACAGCAACTTTGTGGAGCTGCGGTGGGCTCTGCCCAGTTTGAACCTTCTGTCAGCTTTGTTTACACTG
TGAGGGAAAAACCACCTACTTAAGCCTCAGTAATGGCGGACACCCCTCCCCACACCAAGCTAGAGCATCCCAGGTGCGAC
TTCAGGCTGCTATACTGGCAGCAAGAATTTCAAGCCAGTGTATCTTAGCTTGCTGGGCTCTGTGAGTGTGGGATTCACT
GAGCAAGACCCCTTGGCTCCCTGGCTTCAGCCCCCTTCAGGGGATTGAATGGCTGTCTCACTGGTGTTCAGGTGCCA
TTGGGTTATGAAAAAAACTCCTGCAGCTAGCTCAATGCCTGCCCAAACAGCCACCCAGTTTTGTGCTTGAAACCCAG
AGCCCTTGTGGTATAGGCACCCAAGAGAATCTCCTGATCTGTGGCTAGTGAAGACCGTGGGAAAAGCATAGTATCTGGG
CTGGGTAGCTTCGTTCCCTCACGGCACAGTCCCTCATGACTTCCCTTGGGTAGAGGAGGGAGTTCCCGAGCCCCCTTGTGC
TTCCCAGGTGGGGAGACGCCCCACCCCTGCTTCTGCTTGCCCTCTGTGGGCTGCACCCACTGTCTAACCAGGTCCAGTGA
GATGAGCTGGGTACCTCAGTTGGAAATGCTGAAATCACCTGCCTTCTGTGTTGATCTCACTGGGAGCTGCAGGCCGGAG
CTGTTTCTATTTGGCCATCTTTATTTCTTTACAATTGAAAAATAATGATCTTAAATTTTTGTCTCCACAAACAAAAGC
AAGTAAATTATCTCAGTAATATAGTTTTAAGAATAGTTGCTCTTTTTTAAYTGTGGCAGGGGCATAAGGGATAAGAGA
AGAAATAGTTAAGAACACAGGTGGGTCAAATCTACTCTCTGGCACTTACTAGCTGGGGAACTTTGGGCAACTCACCTA
AACCATTTAAGCACTACTAGTTGCCTCTTCTGGAAGATGGCAGTTGTAATAATACATAATTGATAGTGTCAATTATGAGA
TAAATGACATAATGCAAATAAAGTTATTGGCCATGTGCTTGGTACAAAATAAGTACTCAAAAAGTACTTCAAAGTAAT
TTTTTATTCTAGGAGCCATTCCTTACAGGTGGAAAAATTGTGCCTTTTTGCTTACATTTTACTTTGCATTTTCAAGATG
TCTTCAAGAGTATGACCTCATTTTATCCCTTCCACAAATATGAGACATAGAAATTGTTAGCTACATTTTCTAGGGAAAA
AAAACCGAGCGTCAGGAATATTAAGTGACTCAGCCAAGGGTCTACACCTCATAATTAGCAGGATCAAGACCAATGAAAG
TGCTTCTTGGTAAGATCATACCTGAAGCTAACAGACACGTGCAGGCCTCCTACAACATAGGTAGCCATGTTTAGTGTAG
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CAAGCAGATGGCATTGCTTGATTAGAGTCAGGCCATATTTAGAACATTTCTATAAAGCCATTCTCATTTGGGCAAACAT
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AATGACTTTAAGAAATAGATGTATCCTGAAGCAAGATAATATGAGTGTGAGGCTGGACACGTAAAATACAAGGGAAGGT
AGTGGGGAAAAGGCTGAGAAAAGAGTACTGTAGACTTTACATCACAATGCTTAAATTAATCTAAAATGTAAGCTATAAA
ATAGACTAAAATTATATATAGACGTATGTGCGTATGTGTGCATATGCATAAAAGTGTATGTATCTCCATACTTCAAAGG
CTAATATTCACAGATAGATTCAACAATCAAATACGGAGTTAGTTTTGAAGGAAAAGCCAAATAAGACACCTTTGCTAGA
CTTTAAATTTTATTTTAACTGAACTACGGTATAGACATAACCATATAAACTAATCTATTTACTTTGTAAATTAATTG
TTTCATTGTCAATTGAATCAGATGTGAGTGTATTTAGATAAGTCAAGGTAGTAAAAGAGTATAATAAGAAACACTGAGAC
CCAGGGATCATGCTAACTCAGACTTCTGTCTTGAGTCTGCCACATAAGAGCAATATTTCCCTCAGGCAAATTATTTTCAT
AGTTTTGAACCTTGAAATTTCTCATCTGATAAATGGACACACCAATGCCTAACACATAGGGTAGCTGTGAGGATTAAATGA
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TGGTTCTTTCCGAGGGCTGTGAGGAAGGATCTGTTTCAGGTCTTTTTCTTAGCTTGTAGATGGCCTTCTCTCTGTATTT
TCTCTTCGTCTTCTCTCTGTACATGTCTCTGTGTCCACATTTCCCTTCATGTCTCTGTGTCCAAATTTCTCTTTGTCTT
CTCTCTGTACATGTCTCTGTGTCCAAATTTCCCTTTTATTACAAGGACACCAGTCACATTTGAAGTAGGGTGCATCTTAC
TCCTATATGACCTCATCTTAACTAATAACATCTGCAAAAACCCTATTTCCAATAGGGTCCCATTTCTAGGGTACTGAGAG
GTTAGGCTTCAACATTAATTTTGATAGGGGGACACAGTACAACCCATAACAACACTCAGAAGTCACTTTCTCAGACAGA
CCATGCTTGACCACTGATGTAGTTTGGCTGTGTCCCCACCAAATATCCTGAATTGTAGTTCCCATATCCGCACGTGT
AGTGGGAGGGACCAAGTGGGAGGTAATTGAATCATGGTGGCAGTTACACTCATGCTGTTCTCRTGATAGTGAGTGAGTT
CTCATGAGATCTGATGGTTTTTTAAGGGGCTTTCCCTTTTTTGTGTTGGCACTTCTCCTTGCTGCTGCTATGTGAAGAAG
GACATGTTAGCTTCCCCTTCCACCATGGTTGTAAGTTTCTGAGGCTTCCCAGCCATGCTGAACCTGTGAGTCAATTAA
ACCTCTTTCCATTATAAATCACCCAGTGTGAGGTATGTCTTTATTAGTGGGATGAGAACGGACTAATACAACTACCTTT
TCTAAGAACAGTCTTACCCTTTCCTAGCACACACAGTCCCTTGCCCTATATAATATGCCAAAGTAAATCATAGAATCAT
TTATTTTCTTATTCTTTTTCTATCCCCACACTAGTCTGAAGTCCATGGTGGCAGGGCCCTACTTGTCTTGGAGGTTGT
TCTATTCTGGGCTTGGAGAACAGTGCCTGGTACAAAATAGATCCTTAACCTCTGGCTTGAGGTAGGGCTCAATAAGTA
ATAGGTGGTATAAGTATCTAAAAAGAATCTCATTTCTCTATCCAAATTGGTATTTCACTAGCCATATGTGAAAGAACTA
AATTTTCAATTTTATTTTAAATAAATCTAAATTTAAATATCCATTAGTGGCTATTATATTGGACAGCACAGATATAGAAC
ATTTCTGCTACTGCAGGAAGTTCTATGGGACATTTCTAGACATAGAAGCATAGATGAGGATATAATAGTGATAATTGAG
TTCAGACCCGGTGTGCTGCCTGCCAAATTCACAAACACTGCCCTGCATTGCTGCCCTCTCATGACACCACACAGCTCCCAG

Fig. 9.208

TTGATTTGTTCTTCTTTATGGTACCAGGTGTATTGGGTGTATTGCAAGCATTTTCTTCTTCCATTATTAAGTTAAAAAT
ACCCCTTGAAGTTTTGAAAAGCTGTAAAGGGGTTGTCTTGAGAGTCCCATTAACATTTATAGTGAATTCTGTGGCAGCA
ATTTTTTATTTTAATTGTTGTTGTCTTTGCTATTGAGATAAGTAAGCAGTAGGCATRGATAATTAATGTACAGATTCTG
TTTTACACAACCTTTAAGAATGACTCTTTGAGAAACATGTCTTTAAGGACAATGAACTGAGATATCCCTTACTTGAGGAT
AAAATATGGGTGCAGGTACAAGGAGTCAGCAGGTGTCACGTGTGGGCTCATTCTTCCCTCAACCCTGGAAAGTATATA
GATTCTCACCATCCTTGTTTCAGCCTCTGCCTTAGGGATAAGCAGTGCTGGGGTGATTTTATGCTCAGRAAGTAACCAA
GGCTAGCAAACATCTCTCATTGTTGGCAGTAAAAACCATTGCTATGAATAACAATCTATATTTTATAAACTATTTTGG
CTTCACATAAAAAATCTGATGCTTCAAGGAACATACTCCAATGTGCCTTGAAGTATAGCACATACTCCAATGTGTTACTG
AATCCTGAACCACACCCCTAAAGGAGGCATTGATGATCACTGGCTTAACAAAAGGCTTTGGGGTGGGGTGAAGGGA
TAATACATAATGCAGTTAGAAAGTGATAGAAAGCATATTGCTTATATAAAATTCAGTGCTAGGTAAGAATAAAATATGA
AATACATGTGGCAATGAACCTAGAAATATGCTCCAGTTTTGGAAAAAGAATAGTAAGATAAAACCCTAGAATTTGGAACA
TTTTTAAATAACCTCTATTACTATCATTTTAATCTTTGAACAGTGACAAAATATCCACCTGCTTTGTGACTTTTCTGA
AAAGAATACATGGTCACAGGGTATTCTCTATTGGAATACCATTACACCTTCAAACCTTATCACCATCCAGTTAAATTTG
AGGAATGGTAATTGATGACCTTAAGTGAGTCAAGTCACAATTACCTGAAGAATAAAATGCTACTGGAAGGACTAGAC
TCTAAGTTGGTACCTAATAATTCTTATTGCTGAATGTGGTTGTAGAGTAGACCCTTCATTAGGAAGTGAATAGGGGA
AGTATGAAAAATTAATGTGTACAAAATCCAAAGCTTCCTGATAGGTCTTTTGGGCTATTTTGAAGAATAAGACAAACA
ACAGAAGCATTATAATTGGAGGATTATAGAGCCAACAAAGGCAAAGTAGGAGCATATTTATCAGACATAATAAGACAG
AAATTAAGATCTTATTTATGTTTGGAGCATAGGAAGCAATAGAAGCATATATTTGCAAGGTCTAGACATGCCTCTA
TTCATCAATGCATTTTAGTATTTTAGCTCAGCGTTCATTAAAGTTATTACCTTAATTTATTTAATCAAAAATATTGACTC
TATTCTCATACTGCAAATAACTTGTTACATGATTTGGATGTTAATATGATTTTTTTTACCACTGAAAGTTAACTAATGC
TCATCCTTTCACTATTAAAATATTAATGTCTTCTATTTTTAGTTTAGCATTTAAAAAACAAATTTGTAGAGATGGGATCT
CACTGTTATAAAAAAGCTTTCCATTTTTGTAAACTGCAAACCTTGCAAGTTAGCTTAACTAAAATTACATAATAGAATTTT
CAAATTAAGAAATTAAGTCACTCATTTGTTCTGTAAAGCAGTCACCCTTGAATAAAGAAATSCATGGTTCAACATATTT
TTTAAGAATTCAATAGTACTTTGAGCTTTATTGGGATATAACTGACAAATTGAAATGTATATATTTTCAGTGAACAAC
TTGATGTTTTTGATATACATAAACTTTATGAAATGACCATTACCATGGTTCCATATAACCATTTGTGTTTAAACCATCT
AATCTCTTGTTGACTTTTCATTTAAATGTTTGATGTATAGAAACCTCATTTCATTAAATACTTCAGAGGCCAACTGTTA
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AGCTCCGCTCCTGGGTTTCATGCCATTCTCCTGCCTCAGCCACCCAAGTAGCTGGGACTACAGGCGCCCGCCACCACGC
CTGGCCAATTTTTTGTTATTTTTGTTTTTTTAGTAGAGACGGGGTTTCACCGTGTTAGCTAGGATGGTCTCAATCTCCTG
ATCTTGTTGATCCRCCCGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCGCGCCCGAGTCATTGGTTTGT
AATGCCATAAAGAATCCATGAATAATATAGTCGTGTATTTATTTTTGAAAGAGAATGTTCTTTTAATAAAGCATATGTT
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AATTTCCATACACTTAGCAACTTAAACAATATCCTTTTTTTTTTTTAAATGTCGCAGCTCTTTAGGTCAGAAATTCAGG
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GAATCAAGGTCTCTCTCAAGATCATTACTGCCACTGTTGGAAGAATTCAGTTTCTTGCAATTATAGGATAGAAATCCC
TCTTCTTTTGCTAGCTGTTGGCTGTAGCCCCCTGTGACCTTCTAGAGGCCACTCTTGATTCTTATCCCATTGGTCTATC
TTCATCTCAGCCGTGGAAGACCTCCCTCTCACACTTCATCTCTGACTTCTTTTTTTGCTTCTAGCTAAAGAAAGCA
CTCTGCTTTTATAGGTTTTATAGGACTCATGTGATTAGATTAGACCCACCCTGATAATCTTATCTTAAGGTCAGCTGAT
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GTAATAGGAGTCATCTTAGAATTGTGTATTAGCCAAGATGTGGAATTCCTATCAATAAATTTAAGCAGTGTTTCCATTA
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AAAACCTGCCAGAAATAATTCTTAATTAGATGTAGACCAAATGTCTCAATAGTTGGATAACAAGGCAGAGGCCACATAAT
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CGGGGCTGCTTTATATATATTTTTCTCTTCTGTAAAGTGATAAAAAATAGTAAGGCCAAATAATTTATGATAGTTAAC
TGATATGATGCATATAAACTGCTTAGAGTAGCATATACCACATTTATAAATGCACAGAAATGTTAGCTATTTTTTTACTAT
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AATCACTTGTTAAAAATAGCATTGCAGAGAAACATTGAACCTCCTTAAATGTAAATAGCACGATTCATYGAACATCTA
GTCAAGATGACTAATCTTTTTTTCTAGTAACGCTCAACAGTCTAGAAGGTACAGAAGGCGGTGGGTATCAAGGTTTATA
AAAGGTCAGGAATTGGAACCTCTGGGGACCTGGTTGTTCTCATGAGGGCACATAACCAAGATCAGGGTGAGCTAGGAAGTT
TAAGTGATAGGAAGATAATAGATTTATTTAGAGAGGGGAAATTGAGACAGTTTCAAGCCTCGGAGATGGAACAGAACAA
CAGAGCTCCAAGCACTGGGATGCAGGGCTGGAGGTGACCAAGTCCACGACTTTGTCTTTCTAAAGTGAAGTGGAGGTGA

Fig. 9.209

GTCTGAAGAAATTATCACTGTGATTAAGGTTGTTATTGAGGGCAACAGGGCAAGCTAGCTTAGGAAGAAAAAGTTTGGC
CTAGGTRTAAGATGTAGTTGTCTAATATTGAACAGCACAGATTTTAGAACATTAGCATCACCACAGAAAATTCGATTGA
ACAGCACTGTACTAGAGGTACAGGAAAGTGAGAGGGTAGGGTCTTACAGAAAATGTTGGAGCAGAAGAAAC'TATGCCAA
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AAAGCAAAGCCAATGTGATCATCAAGATAATAATAGCCATACATTAAGGTAGTCAAGATCATAGACTTGAGATCTGAAC
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CCAAATGGTTATTGATACCTGCAATAATACTTTTCCCCCTGGAATTATATTTTCTCTTAAATATATGTGTGTATGCA
TACATTCGTATATATAGTGTGTGGGTATGTGTTTGTGTGCATATGTGTGTGCATATATATATATAAAGAAAATAGTGTA
GCGGAAAGAACTCTGTTTTCTAAGTCAAAATATCTGATTTTGAAC'TTGGGGTCCAGTGAGCCACTTAATTTTCAGGTTTA
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TCAGATGACATAAAAGATATGAAAATATAAACTATAACGTATAAATACTGAAC'TGTAAAGTCATACAAATTTAAGTC
ATTAATGTTACTGATTGGAGTTCATTTCTCAGATTTTACATAGTTGTTGGGCACAGAAAATTTGGAAAC'TTTGGCTTTC
ATTCTTGCACAGACTAATTTAGCATCATATTCTTGTCTTTGCTGATTTATCAGGTAAGGAAAAAGTCAAGTAAAATGA
TTTGGATGCCTCACTGCCAGGGAATGCTTGATAAATTAAAGCCTGGCTTCTAGGGACATTCTGACTACTGTTGTTTA
TAACCAAGATCTTTTGCAACCTTAGACTTCCCAGGAAAAGCCTGGGCTTTGTTGCTGCAGGAAGGAGAACAGGGAGGAA
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CTCCTGTGAGAACC'TATAATTCAGGGATGCTAAAGTCACCATGGTGACAAGAAGAACTCAGCTGGCTTCTTATTCAATG
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GCTCCAACCAACCTCAAATCTTAACTCAAATATCCACACTCTAAGCATAGTTTTCTTTCTTTCTAGCTCCTCTTAT
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AATAATTTTGCATCTACTCTCAATTTTGCAC'TTCTTACACCAAATATACTGACATTTT'TAGTGCTAAATCCAGGAAAT
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GCCAGGCACCTAAGTTATTTCAGAAATTCACAGATGATTGTGATATGCTGGTGGTCTTCTTTCACATCCATGCCTTCTC
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CAAAC'TCACTGATGTGTCCAAGAGAGCCTCCTTGCTCCATTCCCTTCCCATCTCATCCAATCTGCTCTACCACCTCCC
CCTTTTCTCCTCCCCCATATTTCAGTCTGAATGAGTGACATCAGATTTTATTAATTTGTACAAGCCAGAACTTGATAA
TCATTTTGAATTGTTTTATTTATCCCTCATGTCCAATCAATTGTCAAGTTATATCAGTTTACCTCTAGTATTTCTCAA
AGTTTTTTATATACATGGGTATTACTTTAGGTTAAGCTAAGTCAGTCTGTGTTGCTATAACAAAATACCTAAAAC'TGGG
TAATTTATAATGAACAGAACTCATTTCTTACAGTCTGGAAGCTGGAAAGTCCAATATCAACGTGCCAGCATCTTGCGA
GGGCTTTCTTGTGTGACATCATATGGCGGAAAGTGAGAGGGCAGGAGAGAGCAAGAAGGGACTGAACTTGCCCTTATA
AGGGCACCAATGCCTCCCATGAGGGTGGAGCCCTCATGACCTAATTACCTCTTAAAGGTCCCACCTCTTAATACTGTTA
CAATGACTATTAAATTTCAATATGAGTTTTGGAAGGGGAGAAACATTCAAACCATAGCAGCCACCATCATCTGTCTTCT
GAAATATTACCACCACCTCCTGACTGTATCCTCTGTGTTTGGCTTTGCTTACTCCCCTAATCTGTTTTTTCACACAATG

Fig. 9.210

CTCAGAACATTTTTTTCTAGAGGATAAATATGATAATATATATTTTCTGCTTAAGACTTTTCAGTGACTATCTGTTGCT
TTCAGAATAGAGGCCAAGCTGTAACATGGCATATATGGCCCATCAGAACCCGACGCCTGTTCTGTTCTTTTGTCTAGGTC
CTCTCTTTCATTCTGCACACAGTTTTTCATCACAATTCTCTGGCGTTCATGATTCTCTCTTGTCTCTGAGCCTTTGTTTC
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GCCTAATTTACCCATAAGCAAAGCAAGGTTTGAATCCACCTATCTAAAATAAGACTTTTAGACATGTTGGAGGATATA
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TCTACCATTTGAGAGCATCGCTAGGGAGCATGACATTATCACAGAGACAGATATATGGTGACTTGGAATTTTTTGTAG
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TTAATGGTTTGAGGGATCATAGATCTGACCACATCAATATTTGGCTCATTTTCAAATAGAAAGTGCTGAAACACCAA
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CCATTTAAGGCTTTTATCTCTGAACCAGAAAGTCATAGTGTACCAGGACTACTCCCAATTCAAACATAACAGGGTTCAA
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GGCGTGGTGGCTCATGCCTGTAATCCCAGCAGTTTGGGAGGTGAGGCCTGCAGATCGCTTGAGCTCAAGAGTTTGAGA
CCAGCCTGAGCAACATGGCAAAACCTGTCTCTACAAAAAATACAAAAATTAGCTGGGTGTGGTAGTGCCTGCTGTAG
TCTCAGCTACTCAGGAGGCTGAGGCCTGAGAATTGCTTGAGCCTGAGAGGTGGAGGTTGCAGAACCGAGACTGTGCCAC
TGCACTCAGGTCTAGATGATAGAGAGAGACCCTACCACCAAAAAAATAAAGCCACATTTTGAATAACTAA
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CATAAAATTCAGTTCTAATTTTTCTCCAAAATAAGATGCTTACTCCAATATAATTATCTTCTTCTTAATTTTACATAT
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CAGGTCAAGTCCAGTCAGGGCAGTGAATAAGAAATGGGTGAGAACTTACTTAAATTTATGCAGTTGCCTGGAAAAAGTT
CAACTTATATAATATATGGGAATATGTGAATGGAGAATGGCTTCCCCCTAATGTGTAAACTGTAACCTCAAAATATTTT
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TCTTCATGGTGAAGCATTACTAAATTAATTTAACGGCAGTCTTAATGTAATGTAATTTTGGTTTACTGATATGTAATCA
GTTAATCTTGAAGATTTTATAAGTTCTGGCATTTGTTCTTTCTCTTGTATTCTCTTATAAATTTGGTCAATTTCTCAA

Fig. 9.211

Fig. 9.212

GCCTCTCTATAGGGCTAACCACAGTATTATAGCTTGTATCCCCAGAGTGAGAGATGAGAGAGACAGATAGGCAGACAG
AGACGGAGACAAAAGACAGAGACATCCCAAGACAGAAGTCAAACACAGTCTTTCATAATCTCTTCTTGAAAGTGACATG
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ATTGCTGTCTAAATCATAGCTTCTTCCAAGTAGCATACCCAGGACAGGGTAAATGCAATTTAAAACTTCTCTCAAGGGG
GTTACTTCATTTAGACTAGTGTGGTGTCTCCTATGCAATTAAATGGTTCTTGTGTCTTATCATTTTTCAATGTCCAAT
ATCCAAGGAAATACATGCACTCATCCACAGAATATTTTCTTTATTTTTAAAGTTTTAATGCTGTTTTTAAGTCCCTTGT
GTTTAGGTGGTTGTTATACTTGAAATGTTAGCAGACATAATACACAGCAAATAAATCTTGATGGTTTTGCTTATCCTC
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CTGCACCTTCTGTCACTTCTCTAAAACGTACAGAACTATTTGGTCAATTGGATTGATTTTTAATGATCTAAATGATCAGA
TGTTTCAACGCTGCACTGCAGCAGCCTGATAATAAATAGGAAAGGGGAGGGGCATGGATGCTGGGCACTCAGTCAGTGC
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AAACAAAGACACCTCTTTATTTTGATTTTGCATATGGGGAACATAAAACAAATGTTTTTGATGAGGGTTTAGGAAGTCA
TCAATAGGGTTAGGAATCAAATTACAGCCTAATTCTACGCGTATCTGTTGACATAAGCAGAACGTCAGTAAAGTGCCTC
CCAATATCTCTAAATAGCCTCTTGGCATCTGAAACGTAGGGAAAAATATTGGTTACTTTTTTAACCAAAGGCCTAAAA
CTTCCAAAGGTATATGAAATCAGTAAATTATAAAACATGTCCCCTTCCCCCTTCTGTAAGAAAGAAATAGAAAACAA
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ACTTTGGTCCAGAGGGTGTAACTAAGGCCCAAACCTGCTTCTGGTGAAAGTGACCCAGAGGTACGCTTTGCTGATAATA
TCAGGAGCATGGTCTTTGTAAATTGGAAGACACTGTCTCCCATCTCCCCATTTGTTATTGCAAGAAAATTAGATTCTCAT
TGAAATAACAGGCCCAAGAATTATAGTAACAGGAGTTTGAACCCCTGTTTGAGAATGTTGGTACTCCTGGGATAAATGA
GATCAATTTTAGCCTATAATAACGGTGCTTTCTACACTTTTTTAGATGTTAGAGAAAGTCTGGCAGCAAAGCAAAGCTTT
GTATATACCATGGGAGGCAGGGTTGGAGAATAAAAGAAGAAAGAGAGACTTAAGAACAAAATTGTGTTTGTATGGGATGT
TTTGATTTTCTTTAAATACTTTAAGTCTCTAGAAATAACAAATTTATTATCTGAAATACAAATAAGAAATACTTATATT
TGGAAAGTTGATAAACTCAATTATTGTGTAAAATGTTTCAAAATTTCCAATGGATAGAAAAAATAAACTTTTGCATT
GCCAATTATTGAATACTTATTTTACCTTTGGAAATAGCAAGTCAGTTACAAATTTTATAAATCTATATAATTGTCTTT
ATACATAATAAAATATTCTGATGTTTTGTGAGCCAATATGCAGTTTTAAGCACCTTCAACCCTGTTACATGTTTACGG
GTTTGGAAAGTTACTGGGGAGGAGGTAGAAAGGTCTGACCTTCTCCCCAGGAATTTAATATCTGGTCAGAGAGAAAGAAA
CAGACTTTTGTGTTGTTTGTGTTGGTCTTATAAACAGTGCCAAAATGAATGTGTGTATGTGTGTGTATGTTTGTGTGCA
CTTGAGCAATCATTTTTGTAGGATAAACTCTTACAAATGTAATTGTTCTGAAATGGTATGTATGGTTTACATTTTTAAA
AATTTGCTAAATTGTACTCCAAAATGTCTCTCCTAATCCATGTTTCTAACAACCTGTGTATGGGTGTACTGTTTCCCACC
CCCTCACAACTGTCATATTTTTAGTCTTTTAAAAAAATCTTAAACGAATGTGATAGGCAAATAACCTCATTTAGTATT
GATTTATTTTTGTTTATCAGTGAGATTGAAGATCTTTTCAAATGGCTTTTGAATTTTATAGTTCCCTTCTGTGATGG
TTATTTGCATCCATGTCCCATTTTTCTTCTGGGTTATTGGTCTTACTCTAATTTATAGACTCTCCTTAAGGCAAATGAT
ACCAAACCATACCATGGCTGTTTACAGTCGACCTCCCAGCTTACGTTGCATTAATAATTGCTTCTTAGAAGCTTATG
TTATTCATCCTTCTCCTATTTCTACACTGCTGTTTTCTTGCTTCCCTAATTTAACATTCTCACCATTGCAGTCAGCAGG
TTAAATGGCAAACATCTGAGGAATCAAGGGAAACAACTATGCATTGAGTACTTGCATGTGCTACATGATTTAATCTATA
CCTTCTAATGTAATCCTGAAGAAATCTAGGAGATATGTATCATTGTTTTCCAGACGAAAAAATTAAGATTTGGTTAAGG
TTGGGAAATGAGACTAAGTAAATAAAGTGGAATAAGAGACTCAGGTCTGTCCGATGATACTAAAGCCTGTGCTCCTT

Fig. 9.213

Fig. 9.214

ATCACTCAAACCTGGGAGGCAGAGGTTGCAGTGAGCCAAGATTGCGCCATTGCACTCCAGCCTGGGCAACAAGAGCTCA
AAAAAAAAAAAAAGAAAAAGCAAAAAACAGCATTTTGTCTATCTACTACATGTCCTCATTTTCAGCAAATATGTTACAATA
TTGATATCCATTTACAAATATTTTAACTACCTTGACAATACCATATCATATTTGTATCTTTCTTGCGTGCTTCCCCTT
TCTTACTCTAATATTTGAGGGGAAAATTATTTATGCCTCCTAGCAAATATGCTTCTGATCTTTAAAACACCTCTGCTCC
AGCAAATGGATCCCAAAGAGATGTGCATTCAAGGTGTGAAAAACAGCAGGTCAGTGGGGGAATGGGAAAGTTAGGAT
TTTTATTTTTTATATTGCTAGAGACAGGGTTTCATTTTGTACCCACTGCAGCCTTGAACCTCCTGGGCTCAAAGGATCC
TCTTGCTCAGCCTTGCAAAGCACCGGGATTACATGTGTGAGCCACTGCAGTGGCTGAAAATTAGTATTAATAGTAATG
TCCTTCTGTCTAGTCCACTGTTTCACTTTCTTTTTCTTCACGGCTTCTCCTTTAGACTCCCTTGGACTGGGAGTTT
AGCACTATCACCTGCACACTAGACCTGCAGTCTATGAAGAGAGGCTGTGAGGGATTGTTGGGCTATCACAGTTGCTCCTC
CCAGAGCAAAAAATATTCAACCCTCCACACACACAGGCAGCAGCCTCATCTCAAATGGACTGTGCCTCTAATCAGTGA
GTGGTTATGGAAGACGAGGAGAAAAGTGCATTTATATTTTCATTAACCTTGTTCCTGCTTCTGTAGCATTTTTCATTTT
GAAAATGAGTTGGGAATTTCAAACATCAATAAAACGTGCTGAGGATTCTGACAACAAAATCCTTTTTTGTGATGTGCTA
TGTTGCCCACTTCTGCTTTTTATTTTTGTCTCCAGTCCCTGTTTGATTGCTAATCTAAAGGGAAAGGGAGGGTCAGTGA
GAAAAGAGGAGAAATCTTAAAAATGAATTTACTATTTGATCAGGTTTGCTACTTTATTTTGTGGAATAATTTAAAGCTC
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CGATTTTTTTAGGAATGGAATATGCCAATTATAGAGAGAGGTAGGCGAGTTAAAGTTTCAAGTGCTCAGCATTATAGA
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CTCTATAAATCTAAAAATATCACAGTATTTCTTTTTTTTCCAAATCTCTCTCTCTCTCTCTCTCTCTCTCTCACCT
CTCCTTATCTGAAACAACTGACATTTAATGAAGAAAACCTGTGAAAAGGGGTTTCGAGAAAATGAGAAGCTAGTATTTG
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ACCAGATACTATTCTAGGCATTAGAGGATATACCAAGGACAAAACAGACCAGAAAACCCCACTACCCCTAGAGCATGTA
CCCCACTACATGCTGATGATAATCTTTAGTCATCTATTTCAAAAATAACTTAAAAATGGAGTGGCAATGTCTCTGCGTA
GTCAATTCAAGATGCCGAAATAAAAATACATGTATATGTGTCTTCTTGTGTGTTTCTTTTTTTTTTTTGTATTTCTCGT
GTTGAACTTGCAATGGTTTTCTAAATGAAATATTTTCATTTTTTAATGTTTCATATATGTGTCTGTATATATATATGAAT
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AACTGCCAATGATGTGTAATTATATATGTGCATGGGTTAGAAGTGGTGGATGGGGCCTTTGGATTTTAAATGGCATT
ACCTGCAGCCTCTGATGCAGCCTCTTCCAGCTTCCCTATTAAAGTGACTGAAAACACTGCAAAAATATATATATATATAA
TTTTTAAAAAGCACTAAAAACCAGTACTGGCAGAATATTAAGAAGCGTTAAATTAGATTGACAATATATATGGTGGAC
CACACATTGTATTCCATTCTAAAAGTAGGAAGAATAATTTCTTAGTTTTACCTGGACCATAACAATACATCTACTATGTC
CTACTACTGTAGTAGTATATGCAGCGATATACTACTTAAGATTTTTTAAAAAGAAATATACAAGCAGACCTAAAGGTGGA
TCAAAACTAAAGAGTTAAGAATAGAAACATCCAGAGGAATACCCTCTTGAGATCATCTACTGGTAAAAATTCATCACAA
GAGTTTTTAAAGAGAATAAATACTTTTGTAAAGCCCCATCTGATTGAACTGCCTTCCCCAGTAAACCTGTGAGAGTGGAGA
GGTTTGAGTATTTTCATGGTAATTTCTTTACCATGTGCCATCTGGCAAATAAAAGAGTTCTTTTCCAGGCAGTACTTTT
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AAACATTCTTGATAAAATTACATCTTTGCTTCAAGCATATTGTAAAGAAAATGGAATCTATTGACATTACAAATAGCAA
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CCAACCCCCACACATGCATAGCCACCCCATTTATCAATATCCCCCCCACAAAAGTGGTACATTTTTTTTTTTTACAATTGAAC
CTACATTGATACATCATTATCTCTAAAACATATAGCTTACATTAGAATTTACTCTTAGTATTGTATATTCTATGGGTTT
GCACAAATTTATATTGATGTATATCTTCCATTTTTGTATCATACGGATAATTCCAACATCTCTGGCATGTCTAGTTCTG
ATGCCTCTTCTGTCTCTTTAACTGTGTTCTTTGCCTTTAACTATGCATTGTAATTTTTTCTTCATAGCTGGGCATGTA
CTGGATAAAAGGAACCTCTGTAAATAGGCCTTAAGTTATATGGGTGGGCCAGGCATGGTGGCTCATGCCTGGTAATCCC
AGCACTTTGGGAGGTGGAAGCAGGTGGATCACCTGAGGTGAGAGGTGAGACCAGCATGGCCAATGTGGTAAACCTC

Fig. 9.215

Fig. 9.216

GAGGTATTCTTACTATCCTATCACCCATAGGTGATGGTTTGACATATGAACAGAGTAGAATAAAATGGACTAAGAAAAT
AATGTCAGATAAATGATGTTTATTACAAAGGAAAATATATTTTTTACATCTTTTTTAAAAAATCTTTGCAATTTGCTACTTT
CTATGGAGACCTACTCTCTTGTATAAGGACTGAATGTTAGTTTTTAAATAATTTAAACCCATCATTATCATCATCATGA
TCAAAGATCACTAACCAGAGTCAAATAATTTGAATTCTCTTCCAGATCTACTGCTCACAAGATGCCTGATAATGAACAA
ATACAATCTTTTTTGGCATCTCCAAAATGGGCATAATAAATGTCTTAATCTCCTCAGCATAATTCTGTGATGCCAAAATA
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GCTACTTAACATCTGAGTGCCTCAATAGCCTCATCTGGAGAACAGGGCTAGTAACAGTCCCTTAACCTCATAGTGCTGTTG
TGTGGATTAAATGGGGTAATACTGTAACCTATCTTAGAATGGGACCTGGGGCATTGTAGATGTTCCCTAGCTCTTAAATAA
TAATATTTAAATGTCTAATATAATATCAAAATTTTAAATTACTTGATTCAAAACATTTCAAAGCTTGTTAAAAACAATGTA
GGCTGAAGTTTTCTGGGCCAGATTACAAATGACCTTATGGAAGAGATTTAGTCCCTTTAGCAAAAAGGGGTCATGAGGA
CACTGCCTAGGCTACAGGAAATCTCAACAAATATTCTCAAGATTCTTTTTATGCTGTAGTGGCATCTTTCTGAACCTACA
AGTTCACATTGGCTTTTAAGGAATCACCCAGTTCTTCTTCTCCTAGGAATCCTCCTTTTAGGATCATCTTTTGATCAA
AATGAAAATTCTCCAAATTATGGTGGTTTTTAAGATTAGTTTTCTTTATACTAGGTTTTGAATTTATGGGACATGCCCT
CCACCCAATCTTGGGTAATATTTTCTGCAATGACAGGACCTCACTGGGGAAATCCTAAATGAAGATAATAGCATGTTAT
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GTAAACAAGGCTTCCCCACCCCGGTGCTTTTTTTTTTAAGGAGTGAAATCCACCAAACCTCTATCATTTGCAAAATATCTC
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TGCAATGTCTGGGAACCCACCTAGAGGGCCCCCTTGACGCCAGCACATCCCCGTGTCAGGCAGCAGTCTCTGCAGCTGAG
CCAGGCTGCTGCAGAGTTAATGTACAGTACCACGGAGCCTGCAAGTGTCTGAGCTGATCAGAGCTGGGGCGGCACAGC
CCAGGGCAGACAAGGCGGCTGCGAGGATTCCAAAGGTTCTGCTGGAAATTCGGCGCTGGGGGACTCCAGCAGGAGCCTG
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GAGTTGGAATTCATCTGTAAAAATCACTACATGTAACGTAGGAGACAAGAAAAATATTAATGACAGAAGATCTGCGAAC
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AAAAGTAACTGTGCAAAACCGAATATTGTCTGAGAAAGTAATGGTTATGCAATAAAAAATACTTTGTTAATATGAAGCA
TCCCAAATAAGTCAAGCATGAGGACTTGAGAACATTTAAATTGCTAATATTTTCATGGAGGAAGAAAAAACTTTGAGA
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TTCCCTGGGGAAGGGCACACATGAATTTCTGATATATCAATTTGTCTGAATTTCTAAAAGAAGGTTAAGGGAACTTAG
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ATCTTTCAGGTAAATGTTCTGGCTGGACTCTGGATGAATAATAGATACCTAAATATAGGTTTCGGAGGGCTTTCCAGCT
GCTTTTATGACAATGTCTCAAATGAAAGCTCCCTGAGAGCTTAAGGTACCACCAAATCACCTGCTGGTTTGTACAGA
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TCTAACAAACACCAGATCCATCCAGAGTCGATGCTGTGGCGTATCTACCTTTTTTGTGCTGACCAGAGCTACTATCCCCAGT
CTCTAGAATGCTTGGGTGACATGCCTGCAACCTCGGTGGCCCACTTCCAACCTGCATCACCCAGAGTTTCTTAGTCAGGG
GGAGCCTTGGTGCCATTGCCTTGCTTGTTCTGTTGGTGAGGGTCAGGCATCAGCAATAAGGTCCTCATTTATTCTTACAG
ACAAATTTACATCAATAGTCTTTAATCTTGAGATTAAAAGATCCTGGAAACAGTTCCTGGCACGTTAGTAGGCATTTGTC

Fig. 9.217

AATTATTTTTCTTCTATGCCTTAGGCTTTTCTTCAGAGTTCATTTTATACCTCTTAAGATTTGCTTGGGAGGGGAAATT
ACCAGTCTCCTTTCTATCAAGTGTACCTTGCTACAAAGCAACAGTTTTTGTCTACCTAAGTTCTGCTGTTTAAGCCCA
TTTGTTTATGTTGTAATACATAGGATCCATGTACTCTTTGAATGCCTGCAATTATAAGCACTTTTTATTTTTATTGCAT
TAGCCTCACCTATACTTTTGACTGGAAAGAAATAAGCTATTCAACTCTGAAGTTTTGGAAAGAATGCAAATTTGCTTAT
TCATGCTCCTCTAGATCTGTAATACATATGTTTGAAAGCTGTATGGAGAAGTTGAGAGTCCTGTTGGTTTTCTTTTGTG
CCTGGAGTTAGGTAACCCCTTCATCTGCTTCACTGCATGTCGTACCAATCTGTTGTTGTTGTTGTTGGCCAAACTAAGCCA
ATGTAGAATGTTAATGCTCTGTATAACTCCTACTCTTCCTGGGCCCTTGCAAGGATTCATTAATATGATGTTGGACTC
AAAGGAAGAAGGTGGTGGTTTTTTTCCAGGTGACCTAATTAATTTTGTGCTTGGTCTTGACTTTCTCAATGGTTGTTG
TTTTCTCTTATTTATAGCCTCCCTTCCAGTACTGCAGTAAGAGATTGTAGGGGTTTGTGACAGAAAACCCCTCTTTC
CCTTTGTCCTACTGTAAGAGCCCCTATAGGGTGAGATTTCAAGGCTCGTGAATTATCGTGCTTAGAATAAAGGTCTCGCC
AAATTGCTCTTTTCATCTCCAAAGACTCCCCCTATCCTCATTCTCACATTTAGAGCCTTTTCTTCGTGAAGGGACCGAT
CAGAAGTTGGCAACAGGCCAGTGCTAAGGAATAATAAACATTGTAAAAGACATATGTGCTTTGGTTTCACGAGCCCTAG
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AGAAGCCTGTGCACAGAGTTCCTACTTGTACTCTCATCCACCTGCCCTAGGGCTGGTGTGTTGATTATTGAGCAATAGT
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TAGATAAGTACTATATAGCTCTTCTTATTTTTCTTCTGGCTAGTTGTTAGAATGGAGAGATAGCCTGGCATTACAGGAA
CAAGTATGGCATGGTTGAAAGAAGGGAAATGCAAGTCAGCTTTCTAGGAATTTAAATTTTCATGTAGCAGCAGTTAAGAG
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GCTTATTTGGTGCTTAGGGTTTAATTTTTAAAGGTAGGACATCTAAATGTTTTAAACTGTCTTTGTGATGCTGAGAGGT
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TACATAATCAGGCTATTGGCTGGAAACATAAATAAACTAAAAAAAATAATTTAAAAAAGCACTCCAGGTTTCTAATT
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GCCATATGTTTTACATAAGTTACATCCTTTAATGGCAATATAACTTTCTGAGAATTTCTTATCCAAATCCCTGTTTGC
AGATGAGGAAGGTAATGTTTACAGAGATTAGATAACTTGCTGAAAGTTCATATATCTATTGCTAGGGAAGCTGACCCAC
TCAAAACCAGGCCTGGCCTGATTTTAGAGCCCAATATATTTTAAATCCACCATATTGTTCTTGTGGCAGATGACTTAGC
ACTCTTGTTTATGGATCTTTTTGAAAATTAGATTAGCAGTAGACCAGAGAAAATAAGCAGATTTACCCTGTTATTTAGC

Fig. 9.218

Fig. 9.219

TCTACGTGTGGCTCTTCGTTCTTTTTTATACATCTTTCACTGTCAAACCTAAGCTCTAAAAATCATCTCCCTAATATCTCT
TCACCTTGCCAAGAAAATGTACTAGCTTTGTATTATATGTAATACAGCATAAAGCTCTGCCTCATCTGGCCTCCTCCTTT
CTAGTCTCATTGTAGAAATGATAATTTTTTATTAATATTATCAGTGTTAATTAATTTCTCTCATTATTCTAGCATAACATA
TCCTATATTTGAGTTTTTTAAATATATCAATTCCTTAGTATAGTGGTTATAATCACACTATATATGTATAGAATGAGGTT
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AAATTAACCTCAATGTTTTTAAAAAGCCATAGTTCTCAGTGTGATGGAGATGGTGAGGATTGGAGAATGTCATTAGGGAA
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TGGGGAAGGAAGAAAAATGTCCTTTGCTAGAAGATATTTATATATATATATATATAAATATATATATATAAATATATATA
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AAGCGGTTCTCCTAGCTCAGCCTCCCGAGTTGCTGGGACTACGCATGCAAGCTGCCACACCCAGCTAATTTTTGTATTT
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CCCAAAGTGCTAGGATTACAGGAGTGAGTCACTGCAGCTGGCCTAGAAGAGCTATTTTGAACGTGCCCTTGGGATAGGA
AGAAAGTCCCTGAGGATCTGACTATAGGTAGGTGCTCAAGTGCTCTAAGGGTGAAAACCTTCTCTGCATAACAGAAGAAG
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GCCCTGGGAAGGCTGAGAACTGCGGGTATACTCCATAAACAGATCCCTCCCCAGTACTGACAGTGGTTCGGTGGGTGAG
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CCTTATTTATTAGAGAAGCAATAATATTTCTCATGGGAATGCATGTATTAATACTTTTCAGAGGTCCAGAAAGTATCAGG
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TTGCCATCTGTACTTGAAGCCAAGGAGCATCAGGTGCTTTGCCTTTCTCACCAGGCTTACGCCATAGCCCTTTTCACCC
CACTACCCACACCTCCCAGTGATGCTGCAACTATTTCTCCTCTGGGCTCCATGCAGTAACTTTCTGGGTTTTTTTTGGTT

Fig. 9.221

GTTGTTTTAGAGCATTTTTGTAGACAGGGTTTTACTCAGTCATCCAGACTGGAGTACAGTGGTGCAGTTATGGCTCACT
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GCATCCATCATTTTTCTATGTTGTGATGTTTGGTATCTTTTCATGTATGTAATGTGCATCTCTTCTTAGGAGTAAGACCA
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CAAATAGCTGGGATTACAGGCATATGCCACCATGCCTGGCTAATTTTTTTGAATTTTGTAGTAGAGACGGGTTTTTGCCA
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AGACTGTCTCCAGAAACAATGAATTTATTGTTGCCTGTTTCTCTGCTAGCTTTTCCCCGTATTTACACAGTTTGCTTTAT

Fig. 9.222

CTGCTGGGATTACCCTTCCCTGACTATGCTCAATCTCTATCTCCCTTGGCAATTCTCCTCCAGGCCCCAGACCTGAGAA
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ACCTTACGGTTCCTGTACAACCAGGTGTATGCCCTCTAGAGCCACCAATAATATTCGAATATTAATATACATAATTTT
ATAAACAGGTGTGATTTTAAATGTCTCTAGACCTTTCCAACCAGAAAAGCTGAATCCAGGTACTGCTGGTCTTTTTCTCT
CCATAGGTTCTCATTTTTCCCTGATGCAGTTATTGCCTTTTTCTGTTTATACTTTCTTCCCTTTAGGTGACTTCACTTACA
AATATAGGAATGCTGATTCATGTTTAGCATTGTGTGGTTTTAGTATTTATTTTCTGTATTTCCACATTGCTATTTTAGC
TCATTCTTTAATTAAATAAAATGCCTTATTCTCTCACAGTATCAGCAACTAAAAGAAGGATACATCACCTACAGTTGTT
AGTGTGAAGTCTTATATTAATCATAGGATATATATTGGTAGACATAAACCTACTGACTAAATTTATTTCTGAAGTATA
CTCTACAAACCATCATTGTATTTTTTACATTTGACAAAGTGGAAGAAAAAAGCTGGCCACAATCTTTAACCTGCCCCA
GTGTCTCTGGCTTTTAGGCACACAGCCAGTCGGCAATGGCGGCAGGGGTGGCATATAAAGATTCTAGATCTCAAAATGC
CAACCTGGCCAGAAAATAAACTCACTATGTGATACCATTTGGAACAAGCTTCTATACCAATGTAGGGGGTGATGGAGGA
TAATATACATTGCAAATTTATGTAAAACCAGACCATGTGGTATTTCCAATAGTTGTTACAACATTGCTTAAAATGATAT
AAATGGCCTTATGGATAAAAGTAGAAGTTTAAATTTTAGAATAGTTCCATTTTAAAGACTAGATTTAGCAAAAATCCTTA
TGGCATATGTTTACTCTTGGGGAACCTTGAGAGATAGTGAACAAGAAACGTTAGATCATAGACTTCTTGGCTGCAAAAA
GTTTTATTTATTTTATCATTTTTTACTCGTATTCCTTTTTATTATTTGAAAAGCTTTGTATAAAGGATTTTTTAGAGATGGCAT
TTCATTAAGGTTATATTTATTAGAAAACAACTTGAATAATTTAAAATTACAAAAGGTAAGTCATTTTATGTATTTGCC
TACTTGTTTTTTGTCTTCTCCCTGCCACCCTCCCTTTCTTCCCTTTGTCTTCCCTTCCCTTCCATCCTTCCATCCTT
TCTTTATTTCATGCCCTGCCTATTTAGAAAAGGATTTGCAGTAAGTTACAAAAGTACAAAAGTCAATAATTTTTTTTTTAA
TTCAGAATGTTAAAAGAGAAAATTGGGTGAAGAGAAAGTGAATATTTCAATAATAAGAGTCATAGTGCTTCCCTCTGAA
GGAGCCCAGAAACCCATCTCATAGTTACGTGGTGGCCAACTTAGCAGGAAAACAAAAGAAATGCTAGACGAGAAGAAC
ACAACATCCATTTCTCAGGAAAGACTAAGTTTTCTCGTACAGAACCCTGAAATGTATTCTCCCTGGGATACTGTTTG
GATAAACAGTGAGTGCTGTAATAGTCAATGTCTTTACCCACTGCTTCACAGCAAAGCAGAATTCCCAAGGTAAAGATGT
GACCATGTAACCTCATTGCCTCACCATTGTTCTCTCCACTCTAATATTTTCAAGTAAATGTGTGTGTCTCTTAGGGTT
GCTGTGCTGAATAAGTGAGAGTCACCTGGCACAGCAGACATCTAGTGTGTGCTAAAGAAATGTACAGAGAATGAGAGAG
AATTTGAATAAAAAATGTAAATAGAATATTATACAGAAGTTAGGAATTAATTCACAAGTGTGTAAGAGATACTCTGTCT
CAGATGTGCAGGCCACTAGTAGTGCAGTGTGACACTCACTGCCTCTCAAGAGAACTTAGTAGTCAGAAAAGTGAAGAAA
TGACATAGCTTTTTTACACTATATTATGTAGAAAGCTTATTTTTTAAATGTTAACCAAGAGCAAGGTCCATAAACTCTAA
TACCTTCCAGAAAGCACAAAAGACATTGAGATACATTTAGAGAAATAGAGGAACTGAGCTAGATATTCACGTGAAATA
GGATCATTCCACATCTTCCAGCAAATAAATTGAGGTGACCAAGGTCCAGGGAGCACAGATGGTCATATTCTAATGAAG

Fig. 9.223

CTCTGGCTCAAAAACACACAAGCTGAAAACAAGGCGAGGATGCTTCATAACAATGTCCTTTTGTAGGAGAAGTGAGGAT
TAGTGATGAGGGTAGGAGTGGAGCAAGAGACAGGGATTAATTTGCATAGCCACCTTGGAAATCCCAAACGTCAATGGTG
TCAGTGATGCTGTTTGCTTGAGTCCCTAAAGGGTTGAATGAAAGAAGTTAACTAGATACAGAGTTCACAGAAGGTCAAT
AATCTTAGCTTTCCAACAAGTTGGAGATCGGGGTAAGGAGAAGGGCAGACCTTAAGAAAAGAGTTATACTTATTGCCAG
GAAACAGTGCTTCCTTTTGCTGTTTTTCCACAAACAGATTTACCTTTGCCCTATGCATTTTCATCTTATTTTAAAAACA
AACTTTTTATTTTAGGTTTCAGGGGTACATATGGAGATTTTTTTTTTAATATAGGTAAGCTCCTGTCATGGGGGTTTGTGT
ACAGATTATTTTCATCACCAGGAACCTAAGCCTACTACCCAATAGTTATTTTCTCTGGTTGTCTCCCTCATACCACCCTC
CACCCTCTGGTAGGCCCCAGTATGCTGATCCTCTTTGTGTCCATGTATTTTCATTATTTAGCTCCCACTTACAAGTGAG
AGCATACAGTATTTGGTTTTGTGTTCTGCGTTAGTTTGCTAAGGATAATGCCTCTAGCTCCATCCACGTTTTGCAGAG
GACATGATTGCATTTTTGTATGACTGTATAGCATTCCATTATGTACCTGTACCACATTTTCTTTATCCAATCTGCCATT
GATGGGCATTTAGGTTGATTCATGTCTTTGCTATGGTGAATAGTGCTACAACGAACATATGTGTGCATGTGCCTTTAT
GGTAGAATTATTTATATTCTTTGGGTATATACCCAGGAATGGGACTGCTGAGTTGAATGGTAGTTCTGTTTTTAGCCT
TTTCAGATGTCACCAAGCTGCTTTCCACAATATTTAGTCTTCTTACACCCCCAACCCCTTGGAAAGTTTAGAATGGAAG
CACATTTCTTCAGATGATTATTAACCTTTGCTTTTTCTGTTTTTACTGCATCATTTAAATTATGGTGTGGAACATCGA
AAAGGGCCCTAAAGCCTCTATTGTCAAGTATGACTGACCTCTCTTGGTAAAGCGAGACCTAGTTCTGACTGAGATTCTC
CCAGTCGTTTCATGGTTAACTTTCCATTTCAGTTTCAGTTGTATTTTCATTGTGTTTCTGCCCAGTGGAGATTCAGCATA
GATGACACCTAAGGAAAAATGCTAGTCATAGTGTGAGAAATCATAGCCTTCTCTTTCAAATGCTTTTATTCATATTCT
TTGGACTTGATTTTAGATATTGCTCGCCAAATTTAGCCATCTTTCTTGGTAGACATTTCTGAAAGTTTAGTGATTAAAA
GCTTAGAGTCTAGAGTTAAATGGAGTCTAGAGTTAAATAGTCTGGGTGTGGACTCTGGCCTGGTCACFTGCCATCTGTG
GGCAGGCCACTTCTTTAAGCCTCAATTTTCTCCCCATAAAAAATGGAGGGGGATGGGGGCCATCATGATACTTACTTTAT
AAAGTGGTTGTGAAGATTCAAGGAGAAGTTGATACATATAAAGTTCTTAGAACAGTGCCTGGCAGGCTGTAAACAATA
TAAAAAGTTAATCATTATTTTTCAGAAAAAAGTTGAGTCAAACCTGAAACAGTGGTACAATGTCAAGAGCAGGAACCTT
GCCTTATTAATTGTGTCTTCTAATCTATGGACTTAGGATCTGAGACTCTAGTTCTTACAACTCATCCATCCTGGGG
CCTGGCATATGGGAGCTATAAACACATCAGCCCTCCTTTGCTCTTTTTCTTAGGGCTGCCCTGAGGAAGAAAGCTAA
TGTGTATAACATCTAGCATTATGCCAGGTATGTGGCCAGAGCTCAAAAACTATAGCTATTATTGTATTATATACTTTG
CTCTATTTTTTATACTCCTGGTTAATGACGGAGAGCTCTGTGAGGGGCTGCTAGAGGGAAGGTTCAATTATTTTAGAAG
CTTTAGTTCTTCCACATTTCACTTTGTAATGTTTGCCTTGCCTTGTATTTATTAACAATAGTTTACTATACTCGTTCC
TAAAFATTAAGTATAGCTTATTATGCCCTGTTAATCTTTCTCAGTGGCATAACAATAATTTAAATATCTGCTTGAAAA
TCTCACCAGCATTTCATTTAATGTGGCTTAGAATGTTATATAATATGAAAAAGTACGTGTTCTTTCAGAAATTAGTTC
CAAAGAACTCCCTATATCTAGAGACAGGCTGTGAGAATGGAGAAAGTTGAGGAGTGTCTCTGTTAGGAATCTTACTACT
GCTTTTAAATTTTCATGGAACACCTATTAAATTGCTTCGCAGCTAGGTTATGAGGAAGGGGAGCCCTGCTCACTCTTTTA
AAGTTAAAGTGAAAAGTTACTTGCGGGTGATTAGGGAAAACCATCTCAGAGGAGTGAGCTGTTGGCATGGAGACCCCA
CATATAACTCTTGGGGGATGGCAAAAACCTTGAGAGAGGATGCTTTATTTCTCACCTCCTAAAATGCCTTTCCAGCTTT
CTGTGGTGAGAGAGAATGAGACAGCAGTCATACCTAACAGTTGGTGAAACAGTTCTATGGGAGTGAGGGAAAGTGAGGG
GCCTAGGATGAGTCAGAAACATGTACATACGGAAAGGGAGAGACAATCCTGATTAAATTGTGCCTTCTTGCAAAAAT
ATATATATTTCCACATTTGATACAAAGTGAAGTATATAACCTTGACAAAGCGATATCTGTGCAAAAAAATTAGAAAA
ATTGCTTTAAATAAACAGTCCTTCAAAGATGTCAGGCACAGAGACTTTCTGAGAAAATGTTTCCAAACTTTTAAGGAA
CAAATAGTATGTTATATAAATTTTCTAGAGTTTGTGAAAAGAAAATCAACTTTTCAATTCATTCTATCAAGTAGACCTA
TTTGCAATAAGAAGGAAGTCCCTCAACCAATCTTACTTTTCGGTAGATAAAATCTTAAATACAACCTTGGTAAATTGAAT
CTTGCAATTTTGTAAAGATTTCATGCTCATGTCAAATAAGGCCGAGTCTGTAAATGTACATATAGTTAATTTTGATCAA
ATCTGGTCCACAGAATCCATCATATTAATATGGTGAAAGAAAAGCAATAGACTTGTATTGACAGGTGCCACAGTGCCAT
TTGACTAAAACCTTAATATCCATTTGTTGTTAGTTTTTTTTTAAAGGATGCAGTGAAATAGCAACAGAAGAATATAGTGT
TTATTTCAATGGTGAAACCAATTTTTTAAGTGTGAGTCGTTTCTATTAAAGTCAGCTTAGTTATAAGCAATAAGACGTAA
CATAATATTTGTATGTAATATTTGTTTGGGAAGTTGTATAGTTATTGTCAAGTTATTGTTCTAGTACAACAACCTTGAAA
CATAAAATATAAAATGAAAACCTGACTTTATAGCTATGTGGGTATTTACCTGGAAGAGCCAAAGGAATTAGTTGAAATGT
TATTAGAAATGATTTCAGTAATTCAGATAGGTAGCCAAATACAAAATTACATATAAGAAAGCTATTAGCTTTTTTTATAGA
AAGTAATAATCAGTTAGATATTATAATGAAAAAATAAAATACCTCTTTATTTACTTACTTTCTTACTGTTGTCCAGGC
TGGAGTACAATGGCACAATCTCAGCTCACTGCAACCTTCACCTCCTAGGCTCAAGCAATTCCTCTGCTTCAGCCTCTTG
AGTAGCTAGGTCTACAGACATCCACCACCACCCCTGGACAGTTTTTCAAATAATTTTTGTAGAGATGGGGTCTTGCTGT
GTTCCCTAGGCTGGTCTCAAACCTCCTGGGCTCAAGCAATCCGCCTGTGTTGACCTCTCAAGGTGCTGGGATTATAGGTG
TGAACCACTGTGCCCGGCAATTTCTTTTAAATAGCACAAAGATTCTAGTCACAAAACATACTACACCTATATTTATGA
GTGAAAATAATAAGAAATTAGAAGGGTTTTTTTTTGGTATTGTTGTGAAGAAAACAAAATGTCTTAACCTAATGCTAAACA
ATCCACAATATTCATTCCACTATAGGCAATTCCAATCACATTTTATTTTTGATTTTAATAAAGTATTTCAAATAATCAAG
CTAACATGGGAAAATCAAGAAAAGTTGAAATAAAGCACAGTGGGTAAAGGTTGGAAGGATGTAGAACACACCATTCCA
GAGATTATATCATAAAAATTACAGTAACATATATCATGTAATATATAATATTGTAAGAATTAGGACGGTGGCTCAAATG
AAGAATTAGATAAAGTTGATTTGGATAGACTATAGATTTCAAATAAGACTCATTTATGTATTTGTGTGTGTGTGTGTGT
GTGTGTGACTGAAGTAGCATTTTATTCAATGAGAAAAGGGATGATTATTCAATAAATGGTGTGAAAGCAACAGCTATC
GTTTGGGAAAATAATAAAGTTAGATTCTTTCTTTAAACCTTATATAAAAATATACATTGGATATATATTTTATTAAA
TATTTATTTAAATAAAGATTTCATTAATATATAATATAATTTTATTAATGTAACAACCTAAAAGTAGTTTAGTTAAA
ATTACTAAAATAAATAAATAAAGTATTTAAATTTCAACATAGGGAGGATATTTCTAAGCATATCACTAGAGCGAGAA
AGCATGAAGAAAATGTCTGATAGATGTGTATACATTAAGGAATCAAATTTTTCTACACATCAAAAACACCATAAAGAAA

Fig. 9.224

ACGTAAAGGCAAATGAAAAGGGTGGGATTCTATTTCCAGTGTTACTAAAGAGACATTGACTTTAATGTCTTAAAAATCC
TTAGAAATCGTCCAGGTGGGGTGGCTCACGCCTATAATCCCAGCACTTTGGGAGGCTGGGGCAGGTGGATCACGAGGTC
AGGAGTTTGAGACCAGCCTGACCAACATGGTGAAACCCCGTCTCTACTAAAAAATACAAAAATTAGCCAGGTGTGGTGG
CACGTGCCTGTAATCCCAGCTACTTAGGAGGCTGAGATAGGAGAATCTCTTGAACCTGGTAGGCGGAGGTTCGAGTAAG
CTGAGTCATGCCACTGCACTTCAGCCTGGGCGACAGAGTGAGACTCTGTCTCAAAAAAAAAAAAAATTCCTTAGAAATTAA
TAAGGAAAAAATGACCTTTCTTCCAGGAAAATGAACACAATCCATAAGAAACACAATTGGTGAGTTACATGAATAAAA
CCCCAATTTTAAAGAAATACAAATTAAACAATAGTGAACATAACATTTTCTCTAAAGAGTATTCAAGATTAAAAACAAA
GTCAAAACTAGATGATAACAAATTCAGCGTATTGACTAAAGTCAGGACATTCTTATTCAATTAGCAATGAAAGTGAAAGT
TTGAGTAGCTTTCTAGAAGAGAATTCAAGTCACACTATGAACCAACAATTTTACTTTGAAGAATTTATCCTAAGAAAAA
AATTAAGAATATGTGAAAAGATTTAGTTACATCAATGTTTATTATAGTGGAACCTGAAAATGTGCCAAGAAAAAAT
GAGCCAAATATTCAGCAGCGGGAATTAGCACAAATATACCTTGATATGTTATCAAGGTAAATGTGTATGATAGGATACTG
ATGCATAAGGATAATTTTTAACATGGTAGATTTTCATTATGTTAAATGAAAAAATAAAAAATACATAAAATGAGAGCCC
ATTTTTATAATAAAAAATATATAGTCATTCAAGGAAAACATGTAGACGGACATCACAAAATACCAACAACAGGTATGGT
TCAAAGTGTTATAGGTGGTTTTTTCAGCACATAGATTACCCCTAGCAAAGAGACCTCTGTGACACTTCGTTTTCTTCATCTG
TGAAATAAAGGAATCATACTAGATTAGTTCTAAATGCTTTTAGCAAAAATATTTTATAATTCTATGAGTATTGATGGTT
AATAACCTTGAGAAAATTGTTATTATTTATTGAACCTTTGGAGTTAGTAGAGTAAACCAAGTTTATCCTGAGCTATCCC
TTGTTTTAGTTAAAATTAGTTAAATTTAACATCCTGATGTTTTATTATAAATTTGTGCATGTGTACATGTGTGTGTTTTGT
GTGTATATATATAGCGCTTTCTCACAAATGCCTGTTTTTGCTGTGGAGTGGGTGGGTGGGTACAGAGTTTCACTGAAAT
GTATATTCCCCTATAGAATTCTTAGAGACACAAATATATCTTGTTAATAGGTTATATATATTTTTTCATGTTTCATCTACC
ATTTGAAACCCTATAACCATTTGAAATCATTCTTGGTTTTTGGTAGGTTTCTTTAATTAACATTCACATTGAGAACTAT
AAGAGATTCAACCCTTGAAAAAGTCCTTTTGCTAAAGAGATTTGCATGTGCTCTAAAAATTGGTGGAATAATGGTTCTGC
TCTAATCAAACGAATGTCACTCTACTCTTATTTCTAGCTAACAGTAAACTGCCTGGAAAATGATAACGTACAACAAAT
ATCTCAAGTGGTTATTTAGTACCTTGTGTGATAGTGAATGAAAAACACACTCAGTTCCTCAGTTTAAATTTGCTGTT
GAACACTTAACATTTACAGTGGCTGGCTGTTTGGAGGCCATATACAATGGTAGAAAAAGTAGCTTTCTAAAAGTGCTT
TGCCTCATTCTTGCTTAACCTGGCAGACGTCATGAACCTTCCAAGAGGGAACATATTTTATTGCTTATACCACCAGAG
GACTTGGCCAATTATTGCTATTTTATTTAATGTTGGTGTGACATAGTGCTTTTGCCTTGAGGAGCTCCAGGTGCTTTAT
AAAATTAGTCAGCAATTCACCTCACAGCCTTTACGCCAGAGAGTGAACATTGAAAATAATAATGATTTCGTGGAACCTAC
AGCCACCCCTAACTCCTATTTCTGTTTGGCCAGTGAAACAGAACAAATATGCCAGTTCCCTCAGAGAGCTAGCAGAAA
AAAGTTCTAGTGGGGTGTCTAGGGAGTTTCTCTCTTGTGTAACCTAGGTGAACTCCTGTAAGGAAAATTTGCCATGCACC
CTGCAGAGCTGAGACTTACTATTCTCTGTTGTAGGACCCATACTTTCCCCAGTAAGGAACCTCCATTTTTTTTTCAGGCCT
TTGCTCAAATATCACTTTCTTAATAAGACCTTTTCATGACCATCTGTTTTTAAAAACAATAATCAAAACCTATCCTCAA
GTGCAGACTTTCTGTATCCTCATTCTGCTTATTTTTCTCTGTAGCACATGTAGCTATTTCAGCACACTTAGATTTTTTA
TTTATTTATTAATTAAGGGTTGACTCCATCCCACCAGATCGTAAGCTTTATGAAGGTAGAACATTTGTCTATTCTGTTG
AGTTATGAATCTTTAGAACCTAGGTGGGTGCTGGTACCTAATGGGTGTGAAATAAGCATTGTGTTGAATGAATGAGCGA
AGGAATGAATGAATAAATAAATTTTCCCCAGTGGAGCAACAAGCAACAGTTGGAGTCCCTTGTGTCAATCAAATATAA
AGATAGCAAAGAGTAAGTAAGCAAGAGTGAACCTGCATCATATCAAGATTTCTTTTATATTCCTTAACATTGAAAATAT
TCATGTGGTGAAATACAGTATTAGAGTTTTCTCCTTTAGTTGCAAATAATAGAACTGAATTTAAGCCGACTTAAACAA
AATAGCAGTTTGCAGGATCTTGAAACTCAAAAATTCAAAGGATACAGCTAGTTTTAATGGTTGTGAGGATTGCTGCAGG
CCCTCTCATGATTTGATTGAGCGTTTGTCTCTGTCTTTCTCTTTCTCTCCTTCTGTCTCACTGTGTTTCTAAGACTTTC
TCTATGTGGTAGCAAAGAAAGCTCAGACAGCACAAAGGCTGCCTGTAGTAATCTTGTGACATGAGAGCATACTCTCCC
AGTAGTTCCAAGGAAAACCCAGGGCTGACTTGAAACGTGTGCACTTGATTACAGGTCAACCTTGAGTCAATTGTGACT
GACTGACCAGGCCTCTGTGTGTCCCGGTTGTGGAGGTGAGTTCCATCAGTAGGAGGGGATTAATCAAAGGAAAATCAG
AATTCCTTTATCAGAATAAGAGGAGAAGATGCATGACAAGTATTCAGAGATACACTGTAACCAGAAAAAATGGTCATATG
TGAAGAGTATCTCATTACAACAGGGAGGAATGTAGTTCTAAGTCTTTTAAAGTTCTGCTTTCTTATAATATCAGATGAC
CACATATTTACAGCATAATTGTGAAAATGATATTTTGAAAGGAAAAAGGAATTCCTATCTGAGATAAACTGTCAAAA
AAGTCAAAAATTTGGCTGTGAAAAAAGTGAACGTTTACTTGAGGTTGTCAATTACTTATTTCTTCTTATATAATTCATAT
AATTCATATATTATTATGACATTCATATAGGCAAGGTGTTAAATTCTAATTTGATTTTATAATTTTTTCAAAGAGACTTA
GAATGTGCTAGAAATCCTAAAAATTTCTGTTCCAATAATTGGTATTTGAATTATGTATATACTGCCCCCTTTTTTTCTTA
TAAAAAATGATAGAAGTTTCATAGATAAATAAAATTCACAGATAATGTTACAAATTTATTTCAAATGTTACTTCTTGCC
TTTTTCTATGCCAAGATACTTAACAAATACTTCAAAATTGAAAGTATTTTTTACTCCTTATTTTATCTATTCCACAGAAT
AGTTAGAGGTAGAGGAAGAATTAGATAATATATTTATTAATAAGTCAACATCCTATTAATGGATGAACATAGAACCAAC
TCATAGGCAACACAGTCAGAAAATCAAGGTGTGGCTTGGTTGGCATGAAAGCATAATGCCCAAGAAGAACTACTGTTGTT
GGTTATGCTGTACTGTATGAGAAGGGCATTCTGACGTTAGCTTTGATACTAGCCTTTTAGTAAGTGAAGTAAAGTTAGT
GAGTATACTCACCCAGTTTCAGATTATTAATTCAGTACTAACTCAGTAATTTGGTGTTACTACATACTGAGATATCCAGC
TGGATTCAAGATTTGCTCAATTCAATTAGCAAATCCAGTGTAAAGGGGAAAAAGAGCTTTGTGGGTAAGCATTCTAA
ATTGGTGTTTTTTATGAAGCCACAATATCTGGTAGTTCGTCCTTTCTCCTTCAGTTCTCTGAAGTTGTAATCATTTCTAAA
TTGTTGCTTTCTAGGTAAAGATGTGAGATAGGGGAAAGGAAGGCTTTCTTAGGAATTTTTTTTCTTTTCATTGCCCAGT
TAAATTGGATTTGTGATTTATATCCATTCATCCTATTTCATACCCATCATATATCTGTATACATCTGTATATATCCATC
CTGTTTCATATCCAATCCAATTCATAAGACTTTTCCATATTTTTTCGTGAGATTAGATTTTTTTTAGTGGGTGAGAAGAAAT
AAAAGGCACATAATATTAACATGCTCCTGGCATTCCAGCAACGAATAATGCCATAGCAATTTTATAAAGATTAAAAATA
CCCTAGCTCACTATGATTTTGATGAGCGCAACGCACCTTTTGGCTTTGTGGGTCCCTCCTCATAAAAAATACAAAATTA

Fig. 9.225

TATTTTGTAGCTGCATTTATGTAAAGATAAATATAATCTAAACTGTTTTGTACTCATTTTTTCAGATTTTAAAAGAGAT
TTAACATTTCTATGAGCTCCTAAAATTATGAGCTCTAGGAATAGTGCTTACTGTGCTATTGGATAAACTAAGCTGGCAT
TGTCATAGCTAAATTGGTTACATTTTAAGCTGTGAATGAAAGGGAATTTGGTAAACCAACAACCTGTTGTATGCTTATT
GTTTATTCCAGTATAAATATATTTTTTGCTATTAACATTAGTTATTACAAGTGGAATACGAAGGGGTGGACTCAGGGTTT
GGAAAGAGAGTTAATGCAAGTGAGTATTTGACTCCATCAACAAAACCTTTACAAGTAAAATCCTCAGGTTTTGTGCCTGT
AATCTTTTTTACAGGATTTAATGCTATTTTACTTCGTCAACAATTTTTTTTTTCTTCAAAGATTTGTATTTTTTAAATGACT
GATGCTATGAAAGAAAATACCCCTCATGATTATATTTATATTGTGAATTTTTTACAGTAGGGGCAAACCTAGAATTTTTTCAT
GCGGAGAGCACTTTGGGTTTTACTCAATATAGCTCTTCAAAGTTCTTCAAAGTTCAAAGTAAAAATTGATGCTAGAGTA
ATTTTCTGTAAATCTAGGTGAAAAGCCACATATCATTTAACAAAGTAAATGCTCATTCATATTTAATACAAACCTCTTT
TTTCTTCTGTTGAAGCTGACTTTTATCCAATTTTTTAGTTTCCAGTTATCTATGGTTTTTGGGAAACAGAGCTAATTTGGA
TTCATGAATTCTGTAAACAAGTATTTCTTGAGAAGCTCCTGGTTGTGAACTACCTAGAATGAACTGGGTAAACTAATA
TCAACATCCATTCTTTTGCCAAACATCACTGATAAACTGCAACATTTTTTTTTTTTACGTGAGTGGTCTTGGAATCAGAC
ATTTCTGCATTGAAAACCCAGCACTGTTGCTGACTGGCTGCATGGCCTGCATCTACTTCTCACATTTCTGACTGCCTAC
CAGGTTCCAGGCACTGTTCTAGGCACTGAAGATGAAAAGATGAATAATATGGAGTTCCCATCTTCAAAGAAGTTACAAT
TTTAAGGAAGACTGGCAAATAAAAGTTACCCCAAAAATAATAAAAATACAGGGGTAAAACCTAGGGCATAGTTATATTT
TGGCTGGTCTCTTTACCTCTGAGGTTCTTTTGCTTTTTTATGTAATTGGATTTTAACTCAGAAGACTGTAAGGTTAAAG
ATGATAATGTATAAAATCAGTACTCTGAGTAATGATCAAAATTTTCTGTTGCCATCCACACATTTGAGTGAGATGTAGT
GAGTGCTAGAAAATAAAGCAGTTCGTATCAAAAAGGGTGTGTTGAAAGAAATAAACTAATCCTTGAGAAACACTGTCAT
GCAATCTCATATCTAGAAATTGTGCCTATGTTCTGTCTATAAATTGATGAATTGGCAGGGTGTACATTTAGGACACAC
ATTGACACTACATATGACATTTGTAAAAACACTTGCTAATTCAGCTAACAGCTCGCTTCTCCCCTTCTCCCCACTACTC
ACTGTGGAGAGAGCTTCCAGCTAATAGCTTAGAATCTCCAGACTAACCAACCATTGATGGAACCAGGTGCTATGTGAT
TGTAGGAATTTATTTGTTGCTCTGAATGTTAACTAAAAGTAATGGGCTTCTTATTTTGTCTTTTCTTTTGTGTTATA
GACGGCAAAGACTGTTTCTCATCAGTCTCATATAGCTAAATTAGATTTATCTCTCATGCTGACTGATTAAAAGGGTTAA
ATTCTGGGCACGTTGTCCCCCTGAGTCTCTGAAATCCATGTCAGAATAGAAGTGCTCATTACCAGACTCGATGAGGAGC
TCCTTTTCCAGAACTTCTCTCTGCTACACTCTAAGGACCAAGCACCTCTGCCTAAATTCGAAGCAACCATGGCCTGTGG
CAGTCTTGGAACCTTTTGCCCTCTGGCATGGGTAGAAGTAGCGTCAATTGGGTAATTCAATCCCTGAGTTCTTCTTCAA
TGAATGAGAGAATCCAGGAATGTGTTGCCAGTTGCCTTTTCTTGTACCTAGTTTTCTTGTCTGCAGAATGATGACAGT
CTATTTCCACACCTCCCCCTGCCACTACAGGATTTATGGTCTGCATGCCCCAGTTGTGAAGCTGGCATGTAGCATTT
TGTTGCCACAGGCCTGTGAGTGTTTAGAAGTTTGTGTTTGTATTTCCAGGGCTACAGTTTGAATATTGAAGGAACAAAC
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CTTGAGCACTGGGAAGGATGCCTAGCCAGAAAAGTCCAGCAGCTGGGTCTGTTAATCTTTTCTGAATGAACCAGCA
GGCAGTTGACCTCTCTAAACCTGTCTTGATTCCTTTATATCTTTGTAGTCACCTTTTTTACCCCCCTCAAGAATTCCCCCT
ATAAAGTTGTTCTTAATAAGACTCTTTTTTGCCCACTTCCCCCTGACCTTCTGAGTTGAATATGAATTTCACTAATTGGT
GAAAGAAACCACCATGACCTTGATAAAGCCATTTCTACTCTGGTTTCTCCCTTCTAAGAATTTGTTTCAATCAAGCAAGT
ATTTGAGTCTAAATGGATCCACTTGGACATTTTAATAATTATCATGAACCTCAGTATCCTTAAAGTGTTGGCTAAAACAG
GCTCAAAAATTCAGATAGTGTAATCTACTTTCTCTCTCCCTCCTCCAGATTTAAGTGGATTAATTCACCTTCACTATGC
CAGCACCTGAGTACAGAATAATTTATTTCTGTGTCGTTTTCATGATGAAAGTCCAGCTGGCATTAGCTCAAATATATGCAG
CTGGCTATTAATTGAAGAAAACCTTCCCTGTCACTCCTCAGTCAAACATGATCTGATTCAGCAGATCCTTTTTCACTCTCA
TCTCTTCTATTTCTACCAGTAGAATAAATGTAAATAACTTTCACTTGGCTCTGCTTGTCTATGGATTGTAAATTACCTGGC
AGCATGCAGACTGACCACATTACTGTTTTTGGAGGCGGCTTGTTTTAGTTGATTAGTATTTCTAGAAGAGTTTTTGCACAT
TTCATCTCTGAGGCCTGAATGATGTCCACTAGCAGGAAATATTGTAGTCTGCCTTTTTTCTTACATACTGTATACTAAA
TTTCAATCTAATTTCAACAAGATGTCAAATTCATAGCACCTGTAGGGCAGGTGTTGGTCTTCTATGAATTGATTATATGT
TTTTCTATCATACTGGATGGTGTACAAGCCAGGTTGTATTTTAAATAGAATTTCAAGAGAGAGATTTGTTCTGGAATT
TAGAAATATATATTATGTGCCAAAATTATTTGATGAGTGCATTTTGTAAATCTTAAATAACAAAACCTACTTGGCATGAA
GAGATGCTATAGTATGTTAACAATTGTGGAACCTTTTTTCAATAAGATGAAATCTGTGCAGAAATGAGGGAGCAGAATCT
TGTGGGAGTGCTAAAATGGTTCATCAATTTGATCCTCACAGCATATGAGAATTCACCAATTATATTACTCTCTGAAAAT
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GCTGCATGACCAGTCATTATTATACGCTCAGCCCTGAAAATGAACTTAAGATGGTAAAATCATACCTTGCCAGGATTTG
GGTGGCTGTGAGGGATTGAGTAGGATTTTAGCAACAGGGACTGGAAAAGAAATCAGAACCTTGGCAGAGAAAAAATAGC
ACTAGGATACTAGAAGAGACCTGGGAGAACTAGCTTGACCTTAGAGGCTGGTCTTGTGAGTCATTTGAGAAATAGTATC
AAAAGGGCATAAGAGGGGCCCGCATAGATTAGGCTATTTGGGGTTACAGGAGATCCAAAGCTAGAAGGGGGCAACAAAC
ATTGTGGTATGAGGCCATCCTAATTTGAACATCAGGCACAGGATCTAGTGTCCATGAAGTAGATCTCAGGGAATGGGCA
GCCTGAGAAGAAGAGTAGATCATGCAGAAGCCAGCAGTCACCCCTAGATCCAGATCTGAGTGTCTATGACCTAGATGCAA
TTAAAATCCTTATAATGGGTCCCACCTGGCAGGATTTCTTTCAGGAACCTCAGGCTGAGCCTGAAGTGAGTAGATTGAGC
TGAAGGGAAGGTAAAGAACTATAAGTTGATAGACTTGCAAAATGGAGAGGCTGCTTTTAGGAAACCATTGATAATAGTA
TCAATCCCATTGATAATATATGTGATTGCCCTTTTATAAGGCCATGTGCTTTCTGCACAATTATGATAAAGTATTTAGC
CAAACCTGTATTCCCTATATAAAGGGGGAGGAGATAGTATCCCATGGGGAATTCCTTCCCATATTATAAACTGGTAGAC
AACTTAAGCAAAAATGTTTTTACTCATATGAACACTGTTGAAGTATAAAGCCAGGTATTTAAGTCATGATAGGTGTTTA

Fig. 9.226

TTTACTTTGAACATTTGCCAGAAGTTTGATGATGGGTATATTTGTAGACTGGGTGTTAGAGGTGTTATATTCTAAAAA
GATATCTTCTGATTAATTTTCTTCATTTATAAAAACAATCTATTTTGATAACATTTAATTTATGAACTTTGGTGTCATG
TTACACTTTATACATTTTGGTTGTACATTCCCTGTAATGCTTACAGAGACAGCCAGTCAGGCAATAATTGCTACATACT
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GAATTCACACTGAGCAACAGGGAAGAGAAGGAAGTGAAGTGTGAGCTGTGTGTGGGCCCTCATTGTTCCAGGA
TTTACAAAGGGCTACGCTGTTCATCATATTAATAAATGGCTCCTGCTCCTTGTCAGCTTAATTGCCACCTTGGCTCC
CGAACAAAGAAATCACAATTGAGGCAAAACATGTTTTGCTATTGAAGGCAAAACCTCCTTAGAGCAATTTTGGGGTGT
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TTGGTACCAAAAAAATGAAAGAAAGAGAGCAAGAGGACAGCATGTCAAATGCCTGATAGAAGTGCATGGACAGAA
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CTAGGAAGTGTGCTGGAGATGCAAATTATCTGGTTCCATCTCAGGCCTAATGAATCAGAACTCTAGGTGGAAGGGAGCA
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GCATACCAATGGTTCATATACCAAGTATATATTTGTCTACTATGTGGAACAGTATTATATTTAAACTGAAGTCTTA
TTTCATGTAGGTTGTACCAAATTTGTGGTGGTAAACATTGTATTTAGGTCATTCAAAAGAGTACAGAAATACAGATTA
GCCATATTTTTCTATAGTGCCATTCAAGTGAATAGGTACACAATACATCTCATATTAAAGAAAAAGAGTCTTTTCAA
AGCCTGTAAGGGTCTGAATAAAGGTCTTAGAAATGTGTCTAATACAGCTGAAATCTAAGAGATTTGGAAAAAGTGGGAGA

Fig. 9.227

AGTTAAATATTGCAAATGCATAGAGCTAAGACTTCTTCATTGAAAGTTGGGTAGTTCCAACATTCATGTGGAAAATCTT
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Fig. 9.228

GACCTTGAGATGTAGTGTCCGTATTAGCCCAGTGGGCCCAATCTAATCACACAAGTCCTTCTAAGCAAAGACTTTTCCC
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TTGCTATTACATCATCAGGCTGCAAATTTCTCAAACCTTTTATGGTGTGCTTCCTTTTGAATTTTCTTCCACCAGATACT
CTAAATCATCTCTCTAATTCACAGTTCACAGATCTCTACGGCAGGGACAAAATGCCACCAGCCTCTTTGCTAAAGCAT
AGCAAGAGTGACCTTTACTCCAGTTTCTAACAAGTTCCTCATCTCCATCTCAGACCAACTCAGCCTGGACGTCATTGTT
CGTATCACTATAAGCATTTTGGTCAAAGCCATTCAACAAGTTCCTCAGACGTCCCAAACCTTCCACATCTTTCTGTCT
TCTGAGCCCTCCAAGTCTCTAGGAAGTTGCACATTTTCCACATTTTCTTCTGAGGCCTCCAAACTGTTCC
AACCTCTGCCTGTTACCCAGTTCCAAAGTCGCTTCCACATTTTCAGGTATCTGCAGTAGTGCCACACTACTCTCAGTAC
CAACTGACTGTACTAATCTGTTCTCACACTGCTATAAAGAACTGCCTGAGACTGGGTAATTTGTAAAGGAAACGGGTTT
AATTGACTTACAGTTCACATGACTGGGGAGGCTTCAGGAACTTACAAACATGGTGGAAGGGGAAGCAAACATGTTCT
TCTTCACATGGCAGCAGGAGAGAGAAGTGCAGAGTGGAGGGGAAAAGCCCCCTTATAAAACCATTAGAGCTCCAGAGAA
GTCACCTACTATTATTAGAACAGCATGGGGGAATCTGTTCCATGATCTAATCACCTCCCATGAGGACTTTCCCCCAAAA
CGTGGGGATTACAATTTGCATTACAATTCAAGTTGAGATTTGGTTGAGGACACAGAACCAGACCATATCAAAGAGTTTG
CTTTGACAAAAGGAAGTGTATCTTTTTTATTACTTATTTACAAAGCAGCTTATTAAAGTTATTAAATAGTTTTCAAAGG
GGCAGCTTTTTTTTTCTACTTTCTGTATTAGTAAGCAACCATCTATGATTGTAATACAACCTGAGGTCTCCCAAGAGAGAA
TGTAGCACAAAACAAGGTGTGCTATCTAAACCTGTAAGAAATGTTGTGTCAGTGAACCAACTCCAGCACAGATATGGAGC
TCTCTCAGAAACAACATAGGAAATTTAGATATGTGAAATTCAAATAGAAATAGAAAACCTCAATTTAGAGTTTAGTTTGC
GTAATATCTTAGAAATGTTTTCATGGTTCAAAGCTGATATTTGACAATTGTGTTAGATCTATAAAAATTCACAAAACAT
CCCTATAATTTTCAGATACAAAATGCTAATAAGGATTTTAAAGTTCAATGTGGACCACAGGGCTTCTGCTTTTGCAGGT
GTACCTTCATTTACATCCTTTTCAGTTGAGGAAGGGTGGGTGTTTGAGAGCAAAATGTATGTATAAGGTAAGAAGAAAAG
AGAAATGAGAGAGAGAAGCAGAAGATAGTGAGCCATCAATAATGATATTAGTTGGCATTCAATTTCAAAGCCAACCACC
CCAATTTAGAGGATTTGCATTTAGGACTAATTAAATTATAAGCTAATTGAGCAGGGACTGAGTTAAGCCTACTGATAGT
GCTTCATAAATATTCATATTAACAATAACAAGATGTTTCCTTTTCAAGGCTAAAAACTTTTTCTAAATGGTGTATACA
ACTTGTGAGTCTTGGTAAGTCAATGTTGTTGCATTCCTGAATTTTTCTATCCCTTTTAAAGGATAATTCTAACTCAAGTG
AACCGAAATTTTCCCTGTAGCAGTAGAGGTCTCTGAAAATTGAGGAAGCTCTCCATGTGTAATGCTCTGAAAATGGCA
GACATTTTCAGAGTCACATTCCTGTATATCATTCATGTGAAATGGCATAGGCAATTTTACTCCTCAAGATTCTTTGCCAG
AATTCGCAATTTAATAAGAACAAGTATTATGAATTGTTGAAGATTCTTCCAGCTCTCTTGGAATAAAGGGTCTTCTCA
AATTGTAGCTTTGGTACATTAATAGTTACTCTGGGGGCCCTAAATGAGTTAGTTAATGTGCAGCTTAAACATCTGAGGCA
CACCCAGAAGTAACTCAGGACTGAGGAATTCACCTTCCCTTGCTACTCAATTGCCGTTTGTGTAAATAGTGGACAGTG
ACACTGTTTGTGTGCAGCTAGCAACTGTCTTAAGTCTTGGGTGTTGTTGGAGCATAAAGTGCACCTCCAGTGCCCTGAGT
ATACCTGTAAGGGTATTTACCATGATTCAATAAGACTTGTTTTTAAATTCCTCTCCAAATAAACACCCCTCTTAAATTTAA
TTTTCTCATATTTCTATGTGGTTATTTATAGTTCAAGAACAAGTATTTAAATATTTAAATGATAGCCATTCAATTAA

Fig. 9.230

Fig. 9.231

TAGTAAACATTACATTTCTTATAATTTACATTAAAAATCTAATACTTCGTTTACAAAAAAGACTTTCAAAGAATATGC
ATGCATTTTATGCAAGGTTAAAGAAGTTTTCAGTGCTGGCTAGTGAAATGGGGGTTAGATGCCCTGTCTATACTACATG
TTTAATCAATGTAATCTTAAATGAGATGACCTTGTAGCCAAGCATTTTTCTTTCCTTTATAAGCAATATTAACCTATA
TTAAATCAAGAAAGCTAATAGTTTTTCGTATTTCTCAATAACCATTAAAGTTCAAACATTGGAAGAGTTTAATATTTTAC
ATGAAAAATCCACAGGCAATAATTTTAAGTGACTTAGAAAAATATTTACCGTATTTTAACCTTAAATGACTGTGTATGTG
TGTGTGTGTGCCTATGTGTGTGCCTATGTATGTGTATGTATGTTTATAATCTAGTAGTTCTCAAGGAGTATTGGAGAAA
GAATATGTGATCACATGTAGTTTATAAAAGCTCTCGTGTTTTTTGAAAGACAAATTTTTGTCAATCTAGTGGTAACAT
GATACTAATTTTTTAATTAATTTTAAATTTCCCTGATAATACATTTAAGCCTTAAAAATATTGTTAGTGATTTCATGTTTC
TTTTTCTATAAAAAATTATTTCTAAGCCTTCCTCAAATTTCTTCCAGACTCTTTTTTCTGATTGATTTTTTAAGAATTTTA
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ATACTTTAGTAGCAGTTTTTTTTCTATGAAAAAATTAGAAGTAGCTTTTCTTGGCAGCGTTTTCTTCTAGAGCAATAGG
GAGGAAAACAGAAAGCTGTAGACTGTTATCTTCGTCCATACATATGTTAGTGCTGGAACCTCAACTCCAAAACTTTGTT
CCTTTGAAAATCATATCCCTCCAGAATTGGTGGAGTGGTGATGGTGTGGTGTGAAGGCCATTGCCAGGACTGAGAAG
GAAAAGCTAAAGAAAGAGGGGAAAAAAATCCAGAGAAAGTGACTGCCTGGGGAGGAGGGATAGGCAAGAGACCAAACAT
CGATTCTAAAGGGAATCTTGCAAATGCTCTGGCAAGTCTATGTATGGGAACATCATTTGCCTAGAAGATGCTCCCCTTT
ACTCCCCTTTTTCCCTATTGGGCTGGTTTAGAGGTCAATAAACTTAATTATTTTACATCTAGTAATGTCCCAGATGATAT
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GTTTTTTATTTACCGTAAAGTTCTTTTACATATGGAATTGAATTCTGTGTGCAGTGTGAGAAGAAATCCACTTTTGTA
TGTTTCCCATATGGCTAACTAGTTATTTCCACACCATCGATTGACTGATCCATCATTTTTTCCACTGACTGGTAGTGTTA
CCTTTGACACGCATCAGTACAGTTGGTCTGCTTTACCTTTCTAGCTATAATTGTGGCGTATCCCTCTATTAATTGCCTC
CTTTAGTGTTATTCAGTGCATTTTATACATTTTGTACGAAGATCTTATACATTTTTTTATTCACATCCACTGTAGACTT
CTGTTGCTATTACAAAATGGCAACAATTTTCAGTTATATTTTCCAAATGCTTGCTGCAGCTATATCCAAATCTAATAAT
TTTTATATCGAACAAATATTTAGAACTTTTCAAACCTTGATGTACTTTTAATGATGTTTCTTCTGATTCTTTTAGGTA
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TTAAAAGGAAGATCTTGGTTCAGTGGTAGGTGAGGCACTGAGAACCTGCATTTCTAACAAGCTCCCAGGTGACAGTGAG
GACACTTGTCCAAAAAAGCAATGCTTTCAAAGACTGTTAAATAGAAATGGAGAGAACAAGCATTCTTTACTTGTGTC
TACAATTAAAGAGCATGCTTCCACATTTTACCATCAAGATTAGATATCTTTGATTAGTTTTTGGTAACTTCCCTCTATTC
CAAATTTGCCATACATTTTTATCGTGAATGGTTGTTTAGTTGAATAATTTTCTGTGGTTTTTCCCCTTTAATCTCTTAA
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TGGCTTCATGTAATCCTTGAGGATTCCTAATGTTTTTAAAGTTTCATTATAGAATGAATTAGGTGTATACCCTCTTATTC
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CCGAGGCGGGTGGATCACAAAATCAGGAGATCGAGACCATCCTGGCTAACACGGGTGAAACCCTGTCTCTACTAAAAATA
CGAAAAATTAGCTGGGCGTGGTGGCGGGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGTGTGAAC
CCGGGAGGCGGAGCATGCAGTGAGCCGAGATCGCACCCTGCACTCCAGCCTGAGCGACAGAGCAAGACTCTGTCTCAA
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AATCTCTCTTACTGAGATTTAAACCTGTATCTTTACAGCGAACTTCAGAAACATGGGAAAGACTGTCATTATCCTTCAT
GAAATCCTGTGGTCCACAATGAAGCGTGTTTATTTTGTACCCTCAGTTTGATTTCTTTTACATTGTTTTCATGGTTTTT
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TCTCTGTGAGCTCTGATGTTAAAGGATTTTGTTCATCCTTACAATGTTTTCTCATTAAGCTCAAATATGCATAATTA
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TGATTCTGTTTTTATATCTCAATATGATACAGAGTTGCAAAATATTTCTAAGCTTTGATCATTCCTTTTGGGCATATTCC
TGTAATAATTTTGAATGATATATTTAGTGGAGAAAATGTAAAAATAATTTAAATCCTTCTAAGAAAAACAAATGT
ATTAATAATTTAAATAGCTGCACTGTAAACCAATATTTCACTATTCAATCATTTGATTACCAAAAAGATAGCGGCATGAAAA
ACCTCTATATTTTTATCTCATGGTTTAAAGATGAATATTTTGCCCTTCATATTTATCATTTTTTTAAAAAAGAACACATTT
AGACTTTGAAAACGATATGTAATGTGCTTCTACATTAATGGGAAAAACAACCTAACTGTATAGTCAAATAAAATATTG
GAATCTTATACCTTGACATATTTTTTTAACTAGACATCATCTAATTTCTTTTCAAATAAGAAATAACTTTTTATCTT
TGTCTTGGTATTACTTTATACTTGTCTGTTTCAGTGGCATTGCTATTCTGTGATAAGATTTTATTACAGAAAATGTCTC
TATCTGTACTTGAACATTAGTCTAATTTTTTTTTTACAAATGTTCCCTTGATTTGACATCTGTCAACTGATTTCTGAGTTAA
TATATCTTTTTCAGTTTCCATTCCTTTGAAATAGGAAATCCAATATTAAACCCTTCAATAAAGATGAACCCTACATCTG
TATATCCAGAATTTTGGTTTGATAAAACCAAACTGATAAGTTCAATGGGGTTAAACATTCCTTGAAGTAAATTTGGAGA

Fig. 9.232

GACATTTGAACACCTTTACCCCAAGTTCATTCATCTATTTCATCTTCTCTTTAATTAAGTCTGTTTGAAAAAGCTTTCA
TTGATTGTTAGCTTCTTGGGAATATGTCCTAAATATCATAAGGCATATGGAGCTCCATGACTCCTCAGAAAAAGAGCACGT
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GGAAACAGAGAAAAATATCTTCATGTCACTTGGCTTAGCCTGCTCCTTTTCAGGGGATAAGTTGCCACAGCATTCAAAAGG
GTGTAGTCTATGAATATTTTTTAGAATCTCTAAAGATGAGGATTTTACAACATGGTATTCAACAGCAGAGTTTAATTTTT
CTTTTTTCTCACTTAACACTGAGTCATTAACAGTTGAGACATTTTACATGACTGTGAGCAGTTGAGCACAAAAACCAGA
TGTTAAGATTGGGTACAGTCAAAAGTTTAGACATCCACACCTGTGTTATGTTTTGTTTATGTTCTGGAGCGCCAAACTT
TGTTACATCTTTGAACCCACTTCAGACCTCAAATTACCACATTTTTTAAAAGCCTCATTAGAATGGTATCATATAGTGC
TGAATCCATAGAACAGAAATTCAGTGTTTTTCAGATTGCTAATCATGTATGAAAGTTTCTGAAATCATATTGTCACACGG
GCTGTAACCTTGGCCTATTCTTCTATCTCCTAAGGAAATTGAATTGACTGGACACTGTTTATTATAAAGATACAGCAGGT
ACTTCTTGTAACTGTGCTCTTCTATATATTTTTGAAAATTATTATTCATTTTCATATTTATCAAAGTGAAAAAGCAA
CATAGCTACTCTACTTATATAGCTCATTTTAAGGAATTTTATACTTGTTTTCTAGTTTTGTGAATTTCTTATGAATTCA
CAAAAATAATGTAAAAGCTAATGGTCTGTCCCTGCTACCATTGTGCATATTAGTCAACAAGTACAAGTAGAATATTTAA
AAGATTTATTAGTTCCTAATTATATATAGAAAAAGACTTTGAAAATTACTTAAGCCATGATTCTACCAAGATGTAACCA
CTTTTATTTTCTTTTTTCTTCCCAATGTTGACTGGTTATATGCATATTAATGGAGCTATAATAATAATTGTATACTTT
TTATCTTACTTTCTAAACCACATTATTATTCCTTAAATATTGCTCTTTCTTCTGACACAGTCTTCTCATGTATCATTTT
GATTACTACATAATATTCTGACTTAATCGAACTTTTTGTATTTTTTGATGTTCAAGCTTCTTTTCTCATTTGTTACTA
TTAGAAATGATGGCTCTAGAAACATCCTTAGAAGTTTTCTTATCTTTTAAATTAACTGCTTTTTATTTTACTAAAATA
ATCTTTAAGAACTGGGATTACTAGGTCAAATTCACAATCATATTTGTGTCCCTGTGCTTTTGTAAACCAAAAAGATGC
TATTTGAGTAAGACCTGCTTGATAACTTACTTATTAATCTTATTTTATTTTGTCTTGCATCACAAATGAGTCAAATCT
ATTTTTCTCTAATTAAATCTTACGATCACCGGTACTTGGATTTAAGTTTTACAGTGAAACAGAAAGAATTTTAAAAAAC
GGTTTAAGTAGGAGAAATTATATTTTCAGATCTCCTTGCTATCTGCTGGAATCACTAGTGCCACCACTTTTCACTGGT
GCTCCTTTTGCTTCACTTAACTAAAGAGTTTTCACTCAGACCCTTATGATTTCTCCAAGATGCATTACATTTTCTGCT
GTTTCCATATACCTGGGAAATAGTTCTTTTCACTTACATCTTCTGTCTCTATATTTTTTCCAGGTAAGTCATGCCTCCTTT
AAATTTTTTACTATTTTTTATCTATGTCTGTCTATCTCTACTAAGATGTTCTCTACCTTTGTTTCTGTCTACCTAAGTAG
CTAGCTACCTACCTACCTATCTGGAACCTCTACCTCTGTCTCTTATACTATGCCCATTTCTTCTCTTTTCTCTTTAAAT
TTTTTGTTGCTGTTTTCTCTTTTCTCTATTTCTGTATCTCGTTTGTGTCAAATTGCCAGACTACTTAATTAGTATATT
CATGTTTGTTTTTGAATTTCTATTACTGTATTTTCTCTTATTAGTATTTGTTTTTATAATTATTATTTTCTTTACTTT
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CCCTTTGATTTCTTCTCTCACTCAAGTGTTATTTAGAGTACTTTTTAAATGTTTTCCAAAGGAATGAGTTTTTAAATA
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TTCAAAGTATAATTATATATAAGTTTATATAATTTATAAAGTTTATAATATAAACCTATATTTTATATATTTTATAACT
TAATATAAAATTATAAAGTTTATAATATAATTATAATATTTCAAAAATATTAATTTAGGCTTCTTATTTATCCTGCTTTC
TAATGTTTGTTTTTCTTTTTCACGTCTTTTCTGGCATTATTTTAGATCAAAGTGATTTCTCCTTTTCTCATTGTATTTT
TCCTTCTAGTTTGTAAGTTTTAACTCTAATTTCTATTATTTTAAATAATAACCCTAAAAAATTTTCCAAATATGTAACCT
AAGGCTGAAGTTAAATAATATCTTTACCCTTTTACTGTCTCCTGAGAAAATACAAAAGAATAGATTTTTGTCTCTTC
ATATTTGTATTACATTGTAGACATTATCATTATTATTATAGTTGTAGTAAGTCATCTTTGGTGAACATTTTCTAACACA
TTTTCCATTTTCTTTGCTCAGCATTTTTTTTTTTTTTGCATCTGAGACCATCCTTTTTTATTTAAAAATGTCTTTTAGAA
GTTTCTCTAGTGACTCTCCAATAGTAGGGAATTCTCTTCATGTGTGGTTTTCTTACTATCTTATTTTGTGCTTAGACTT
GGAAGGTAGTTTTGTGTGTATGCATTACTGGGTGACAACACTTCTCTTCACCTCTTTAAATACGATACTACCTTGT
CTTTTGAGAAATCACTTTTCAGTCTGACCAGTGTTACTTTTTTCTTTTTTGGGTGCTTTGAAATCTTATTTTTATCTTT
GTTGTTGTGCAATTTCACTCTAATGAATCCAGAAGTTGAGTTTTTCAAAGTTTACTTAAATGTATTGGTATTCCTAA
ATATGAGTGTGGTGTCTTTTATTTATTTTGGAAATTCCTAGCTATTATTGTTTCAAATATTGCCTCGTTGCCATTCC
TTTTAATCTATCCCAACCCTAGTAAGACTTATGTTGGCCCCACTCTGTCACTCACCTATCTTAACCTTGCTTTTCATATT
TTCCACTTCTTTTTTCTACTGCATTCTGAATGTTTCATTCAATATATGTTTTCTGGCTCACTAATTCAACCTTCAGTAGAG
TCTAATTTGTGTTTTCACCTGTTTGTGAGTTTCTAATTTCAAGTTATTAGACATTCAATTTTAGAAGTTCTATTGGTCT
TTATTCTAAATATGTGTGTTGATTTTTATTGGTCTTTTGCTCATTTACTAAATTTTTTATCATTTCTTTAAACAGCTTA
AATATTTTTTTTTTATTTTAGTTCCCTTGATAGTTCCAAATATCTGTAATCTTGTTGGTCTACCTGTGCAGTTTATTCCACT
GATTGTCTCTCAGGCAAGTTTATTTTTTTTTATGCATTTTCTATATTTTTTTTACATATGTGAGCTCATACTCTTTAGTAT
TTTATCTCTGGAAATTATCTGTGTTGAAAGTGATTTTTTCGAAAGAGGATTTGCTTATAATTTAAGTAACCTGCATGTCT
GCTAACTATAGATTTACTTTAAATGAAATGTTCAACTTTTGGGCCACGCAAATATCCAATGCCATATTACATATCGCAA
GTATACACTTGTGATTAGAAATTTTTGAAAGAAATATTTGTTTTAAATATGCTGCCTGAAACCAAGACAAGTCTTGTTCT
TGCATGAAAGATTGTAGCATGCCTAGTTATAGGTATTGACTTTTGGGTATAAGACTTTCTGTGAAGTCTGAATGTGAGT
TCTTTCTCTGTTTAGGCCCTTGTTTAGATTCTTGCCCTGGAACCCAAAGAATTGGTAAATGCCTTCAGGAAAAAACA
AAAAACAAACAAAGCAAAAAACAATTCTAGTACTTAGTTATCCCTGTGGGACCAAGCTTTCTTAATTTCAAGCCTCAA
AACTTTTAATTTAATGATTTAAAGCAATCATCCAGTATAATTTTAGAGCAAATTTTGTGGAATTATACCATAAGAGGA
AAAAGCACACAAACCTAAATGTAAATGTGGTAAATTTCCATGAAGATAACATACCCATGTAAGCAGCAATTAGATTAA
TAGGCAGTATATTGCTAAGCTGTAGAAGCCCCCATTTGTGTACCTTTTCTGAGATTCTGCCCCACCCAGAGTAACCTACTA
TTGTGATTTCTAAGATAATATTCATATTATTTACCAAAGAGAGTGTGTTCTTTGTTCTCTAGCTTCTTTTCTCAACAC

Fig. 9.233

TGTATTTGGAAAAGTTATCTTTCTTTTTCTTTCTTTTTTTTTTTTTTTTTTTTTTTGAGACAGGGTATTGCTCTGT
CACCCAGGCTGTAGTACAGTACCATAGTCATAACTCATTGCAGCCTCAAACCTCTGGGCTCAAGCAGTCCTCCTGCCTC
AGCCTCCCAAGTAGCTAGGACTACAGACATATGACACCACACCTGGCTAATTTTATTTTATTTTTTTGTATAGACTGGG
TCTCGCTTTGTTGCCAGGCTAGACTCAAACCTCTGGGCTAAAGGAACATTTGTCTTTATATATGTAGTTGTAGTTCA
TTTTCATCTTCATACAAATATGTTATAAATTATTTATTCATTTTATTGTTTTGTTTTGTTTTGTTTTGTTTTGTTTTGAG
AAAGAGTTCCACTGCTTTGCCAGGCTGGAGTGCAGTGGCACGAACCTCGTCTCACTGCAGCCTCTGCCTCCAGGGTTCA
AGCAATTCTCCTGCCTCAGCCTCCTAAGTAGTTGGGTCTACAGGCACCTGCCACTACACACGGCTAATTTTTGTATTTT
TAGTAGAGATGGGGTTTTTGCCATTTTGGCCGGGCTGGTCACGAACCTCCTCATCTCAAATGATCCACCTGCCTCAGTCTG
CCAAAGTGCTGGGATTACAGGTGTGAGCCACTGTGCCTGGCCCATTTTATTATTGGTAAATATTTGTTTTATTTTTATT
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CTGTGCATTAATTTTTTTTTCTGGACTTTTGAAGTGTCTGTATGGGAACTATTCTCCATTTAGCTAGCTATTTTTGC
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TACCTGTTGATTATGCATAATTTTCTTTAAATGATGGATTCTTAGCCAATAAGCTATTTTAAGCAGAAGCATTTTTTTT
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AGTATGTTACATATGGGAAATCTGTATACCTTCATAATTTCACTGTGAATCTCACTCCTCTAAAAAATGAAGTCTTGAA
AAGTTAGTATTTTAAATACTTACCTTACTTGTAAATGTGAACCATTTATATTAATTTAGCAAATCAATGTTTCAAATT
ATTAACAACCTAGACAAACTACTGAGCCATATGTTAGGTATATTGTAAGCATGTTGAAGTCAACAACATGACTTCTACAC
TTTCTACTTTCTCACGGTTCCCAAGATATCTATTTTGACATAATTATTGAAAAAAATTAAGCATTATTTCCATTGT
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GAAGTCTTAGAATGAGTTCATTAGAATATAGACATTGAATATGAGGTAAGTCTGAGTTGAAATTTCTATCTCTGACAACCT
GCTGGATCTGTTTTTGGTTTTCCAATAGTAAATGGCACTAATAATGATATATGCAAAATGTAAATGCTTAGCACAGTG
CTTGGCACATAGTAGATTCTAAGCTGTAGTTTAAACAGCTATTGTGCTATGAATAAGCCATACGTTGACATTTTCCCTC
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GCTGCAGTTCCCTTAGGTTAGCTTCTATGCATAACCTGGGAATGCATATTTTAGCTTTTTACTTCCCTCCCAAAGAAAG
GAGCAGAAGAGGAGAACCAATAGCCAAGATCTAGAAGATGAACTGTTGTATCCATTGTGTGGATTGAGATAAGCCTC
CACCCAAAATATGGTTCAAATGTTAAGACTAATGAGCAGCCCTACACTTCTCTCTCTCACTTTTACACACACACATA
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ACAAATGCAAAATGAATTGTGGCTTAATGTGCATATGTATATAACAGATACTTTGTTCAACTATATTATGAATACTAT
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CATAGGCCAAGTGTTAATTTCTGGTACACATGACATTTAGCACATGCTATTACATTTATTGATTGCATATTTTTATCTGT
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CATCCTAGACCCCAATATATAAAGTAAATGACATTTTGTCTTGTCTTAATAAGTTCACAGTATTTTCAAGAAACATGGG
AATACAGAATAACAGATAAGAAATATTATAAGACTTCTAGTAGAGAAATATGCTTAAATCATATGTATGTTTCTTTAC
ACCTAAGAACCTTTAGATATAAATGCATTTTAACTGTTCTGGAGAGTTCCAACCTCCAAAATAGCCTTTGTTTCATATCAA
GAAAGTAAACCACCTTCTATTTTCAAGGTGGGCTTACTCACCATACTGGGGCCACATTACTGTCCAGCTCACTGAAAGG
TCATAATCCAGGATGAAACCAAACTTGAAATTTATAGTGAAACACAGTAGAATAATTTAGAAGCATATACTTTGATGTT
TTTAGAAAGTAAGGAAATAAACTTTAATTGAACCTTGAATAAACTCAGTTCTGAGCATTCATTCTACTCTGCAGTTG
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CCCTGGGCAACAGCATCACCCTGTACTGTGTTTCTTGGTTGTCTATGGGAACCTTTGCTGTGAACCAGCAGTGAGAG
CACTTCCATTGAATAAAGCTGCCTCTGGATAGCCATGAGTTGCATGAAGTTATTTTACTAATTTTATTTGCCTTTTACT
TGAAAGCATAAGTTTCTGAGGATGTAATTACAGTTTCCCTATATTTCTACAGAAGTAGTTATAGATGATGGACCTCCT
GAACTTTTATCACTTTGCCATACTCTGTAAAATTACCTAAGAGCTCCAGAGCATGAAATTAACACTCAGGAATATTGTA
GCCTAACCTTTTTTGTACTCTGAATAATACTGAATTGAAGGGTCTTCCACAGCAAGCCCTCTTTAAATTATGCTTTCT
GATGCTTTACCTAAGGCTATACACTTGTCTTTAGATTCTTCTAGATTATTTTCTCCTAATACAGAACTTAGCATTTAAA
TGACAATTTTAAAAATTGAGATCAACTCTTCAAAGAAATCAAGTAAAGCAATTTTCAGCAAAAAAAGCCT

Fig. 9.234

TCTAATCTCCTACAAAACACCTCTATCCTATCAAGATATTGTGATTGAGAATTAGAAATAGTGTGGAATCAGATGTTAG
TTTGTCTTTTCTTTTCTACTGGAATTTCTCCATGCTTTAATTTTTATATCAAGAAAAACAATTCTTGCAATATTCCTTA
AGGAGAGTTTTACAAAAAAGAGATAAAACAACTACACACACACACACACACACACACACACACACACACACTA
CCTTAAGGAATAAAGAAATTATCTGAGTGGAATAATGGTGCTTTCAATTTTCTTAATTGAATTAAAATAATTATCTA
AGTCTTTTTTGATCTTGTCCTCAAAGTGAATAAAATGTCTTTTTTTAAATTTATGTTTCATTAGATATATCTCCATCTTTT
CAGTATTCACCCCAAGTTTTAATTGGGTAGAAGAATATGGAAGAAATTCCTATTGCTCAGACACCATGTTTAAAGCTTT
CTGAATGTGATCTAATACCTCCATTGGTTGGAGGTAATTAAACAGAGATCTGTTCTTTTAAACAACCTATTTGAGTGCTT
TAAATCTGAAAGTCATGTCATAAATGTCAGACCTCTTTCTCTTTGCTTTAATCATCTCAGAAAGACTTATCTGTTAATG
GAACATCTGCATATAGGTATTTTGTGACCACAGGTGTTCCACATAAGTTTAAATCTACCTATTAGTTATATGTTTCATGC
TATTGAGGACCTTTCAATAATAAACTTTACTGAGTGCTTACTCTCTTCTAAGAGTAATGCACTATGCTAGGAGACAGGA
GGGATATAAAGGTAAAGAAACCATGGCATGAATCTTTAGGGAGATTATAATTTAGTGAGAACACTAGTGAAAGGAGGC
AAAGGAATTAATAGAACAAGACAGTTTTCTCTGAGTCTCAAAGGCACTGTAAAAATTCCTGGGGAACTGCAAGCAAGA
GCAAAATGTTAGCTATCATAAACATTCTCATGCATCTTCAGGTTTTATCTTCATCCATAAATTTACCCATTAAATTTAG
ACATTCTACTATCTATTCCACTTGACATTCCAATTAGCAGCTCAGCCTCAGCCTATCAATATGAATGCACATCGCTTCC
CTGTTCTCTTGCTGTGCCAGTCTTTCCCCAGTGTAAGTTAGGTAGTGCTCCAATCAGTCACCATCACCATACTGTGCTG
ATTTTATATTCTGAATATTTTTTAAGCTTTTTCTTTTTATCAATTGACTACACTAGTTTAGATCTTCATCATATTATT
GTACACAACCTGCTGGGTTTTTTCTGTCTGGAAAGTCTATTTTTCTCTTTCTCCAGCTGCCTCTTTCTTCTGCCTCT
GGATGCATTTCTCATTTTTCTGCCTTCCAAGATGTATTTAATACATTCCTCTGTGCTCTCATATGCTGATACATGTCT
GTCGCTGTATTTGCTGTGTAATATCCAAATTATTTGTGCATGTCTTTGTTTTCTCCAAAGTTGCAAGCTCCTTAAGG
TCAGGAGCTGTTCTTTAATATATCTACTCTACCTAGTGTAAGTCATGCTTAATATTTGTTGAAGTAAACAAATCTGAACC
CCAAATGTATTATGTAATCTCCATATCCCAGCACAGATAAAGGATATGCTCTTGAATTGTTTTATTGGGAGAAAACAGC
TGACAGATTAGGCTACACAACAACCTAAAACTAAGAAATGACTACATGATAAATTGTAATGAAAATCGGAAGGCAGACT
GACTGAGGAATATATCTCTAACAAGTATTCAAAGATATAATTTTGTATTGTTTAAAAAGAATACAATGAATGGAGAGAG
GGTTTGGATTTACATTCAGGACTTCTGAAGACCAAGTATAAGAGATTCAAAGTGATAATGTGAATCTGATTAGAAAA
TTCAATTTTGGAATTTAAATAAGAAAAAAATTAGTTGGTCTAGCAGGTGAGGCTATAATATGGAAAAATGAGAAAGGGA
TACAGCACTGAAAGTGAAAATGTGGAAAGAGAGAAGATCAGAAGGGGAGAAAAAAGAAGAAATAAAGGGTTAGAGTTT
ATGAAGTGTACCAAAAGAAATGATACAATGGCCAGTCTGTGAGGGTAGTAGAGGGATCTCTTTATTGTATTTTCTTAT
TTTTATTTTTATTTTAGATTACAGGGAGTACACGTGCAGGCTTGTTACATGGGTAAATTGCGTGAAGCTGAGGTTTGGGCT
TGTAATGATCCCAGTGCACGTAAGTACATAGGACCATAGGGAGTTTTTCAACACTTGCTCACCTCCCTCCTCCCACC
CTTTTGGAAATCCTCAGTTGTTCAAGTGTTCATATTTGTGTCCGTGTTGCCTCCAGTGTTTAGCTCCCCTTATAAGTA
AGAACATGTGGTATTTGGTTTTCTGTTTCTGTTTCAATTCAGTGAATAATGGCCTCCAGTTCCATCCATGTTGCTGC
AAGGAACATGATTTCACTCTTTTTATGGCTGTATAGTATTTCTGGTATATATGTAGCACATTTTCTTTATCCAGCCCAC
CATTGGTGGGCATATGGGTTGATTCCATGTCATTGCTCTTGTTGAATAGTGCTGTGATTACATATAAGTGCAGGTGTCT
TTTTGGTAGAATGGTTTATTTTCTTTGGGTATACACTCAGCAGTGAGATTGCAGGGTCAAATGGTGGTTCTATTATTA
GTTCTTGAGAAATCTCCAACTGTTTTCCACAGTGGCTGAACCTAATTTGCATTCCTGCCAACAGTGTTGCAAGCGTTCCC
TTTTCTCCACAGTCTCACCAACATCTGTTACTTTTTTGATTTTTTTAATATACTAGCCATTCTGAGTGGTGTGAGATAGT
ATCTCATTTGGTTGAGTGTGGTTTGCTAATATTTTGTGGAGGACTTTTGCATCTGTTTCATCAGAGAGATATTGGCATGTA
GGGTTTTTGTGTTGTGTCTTTGCCAGATTTTGGTATCAGAAAGATACTGGTTTTGAATTAGGAGTTCCTCCTCCTTGA
TTTTTTTTTTTTTTTGAATACTTTCAAGTAGTATCTGTATCAGCTCTTATTTGTATGTCTGGTAGAATTCAGCTGTGAAT
CCATTTGGTTCAGGGCTATTTTGGACATTAGGTTTTTTTATTACTGATTCAATTTCACTTGTATTGGTCTGTTCA
GGATTTCAAGTTTCTTCTGTTTTAATCTTGGGAGGTTGTGTGTGTCAGAAATTTATGCATCTCCTCTAGGTTTTCTAG
TTTTTGTGCATAGAGATGTACATAGTAGTCTCTGAGGATCTTTTATGTTTCTATGGGATTGGTTGTGATGTCACTTTTG
TCATTTCTGATTGTGCATATTCAGATCTTCTCTCTTTTCTTGTAAATCTAGCTAGCAGTATATCAATCTTTTTTTATCC
TTTCCAAGAACAACTTCTCATTTTGTATATCCTTTGTATAGTTGTTATGGGTCTCAATTTCAATTAATTCTGTTCTGA
TTTTAGTTATTTCTTTTTTTTTCTGCTAGCTCTAGGTTTAGTTTTGTTCTTGTTTTTCTGTTTTTTTTTTTTTTTAGGT
GCGATTTTAGGTTGTTAATTTCAAGATGTATCTTCTTGATGTAGACATTCAGTGCTAGAACTTTCTCTTAACGCTGCC
ATTGCTGTACTCTAGAGGTTTTGGTATGTTGTATCTCTATTTTTGTTTGTTCAAATAACAGTTTGACTTCTGCCTTAA
TTTTTTGTTTATTTTAGAGTCATTCAGAAACAAGTTGATTAGTTTCTCTGTATTTGTGTGGTTTTAAGAATTCCTCTTGC
TGTTGATTTCTGTTTTTTTTTCCACTGTGGTCTGAGAAGATGCTTGGTATGATTCTGATTTTTTGAAATGTATTAAGACT
TGCTTTATGACTATGTGTTCAATCTTGGAATATATATCATGTGCAGATGAGAAGAATGCATATTCTGTAGTTGTTGGGT
GGAGTATTCTATAGATGTCTATTAGGTCCAATTGGCCAACCTATCAAATTTATTAAATACAAAATTTTTTGTAGTTTTT
TGCTGTCTATTGCTTTCACTCGTGTGCTGAAGTTCCCCACTATTATTGGGTGGCCATGAAAGTCTTTTTCATAGGTCTA
TAAGCAATTATTTTATAAATATGTATCCTCCAATGTTGAACATATATATATTTAGGATAGTTAAGTCTTCTTATTGAAT
TGAACACTTTATCATTTATGTAATGCCTTTCTTGGTCTTTTTTTACTGTTGTTGTTTAAATGTCTATTTTTTTCTGATATTA
GAATAGTGATCCTTGTTCTTTTTTTTTGTTTCTTATTGTCATGATAGATCTTCTCTATCCATTTACTTGCAGCCTAGCCT
ATGGGTGCCATTACAAGTGAAATGGGTATCTTGAAGTTAGACTTGTTTTTAGTCTAATTTGCCACTTTTTTGCCTTTTAA
ATGGAGTGTTAAACACTATGACAGGAATAAAACCTTACATATCAATATTAACCTTTGCATGGCCAAAGTTAATATTTTA
TGTTAAACACTATGGCAAGAATAAAACCTCACATATCAATATTAATTTTGACTATGCAAAAGTTAATATTGATATGTG
AGGTTTTATTCTTGCCATAGTGTTATTATCTGGTTGCTTTGTAGTATTGATTCAGTCACTGCTTAGGGCCTGTGGGCTA
TCTGCTTGCAATGTGCTTTCATGGTAACAAGGATCATCCTTTCTTTGTTTTCATGTTTAGAACTCTCTTAAGTATCTC
TTGTAGGTCTAGTGTTGGTGGTGTGATGGATTCTTTTAGCAATTGCTTATCTGAGAAAGAATTTATTTCTCTTTCATTTATG

Fig. 9.235

AAGTATAGTTGGTGGGATATGAAATTATTGGTTGGCATTCTTTCTTTAAAAATGCTAAAAATAGTCCCCCAATCTCT
TCTGGCTTGTATGGTTTCTGCTGAGAAGCCTGCTGTTAGTCTGATGGGTTTCCCTTTATAGATGATATGACTGTTTTCT
CTAGCTGCCGTTAAGTTTTTTTCTTTACGTTGACCTTGGATAGTCTGATGATTGTGTGCCTTGAAGATGGTCATATTA
TATAGTATTCTTCCAGGAATTCTCTGGATTCTTGTATGTGCATGTTGACTTCTCTGGCAAGATTGAGGAAATTTCCCT
GAATTATATCCTCAAATATGTGTTCCAAGTTGCTTAGTTTCTCTTATCTCAGAAATGCCAATGTCACCTTTACATAACCC
CATATTTATTGAAGGTTTTATTAATTGTTTTAAATTATTTTTTTCTTTATTTTTGCCTGCCTGGGTGATTTCAGAAAGC
TAGTCTTCGAGCCCTGAAATTCATTCTTCTGCTTGGTATAGTCTGTTGTTAAGGCTTCCAACGTATTTTTGAAATTCCC
ATAGTTAATTTTTCAATTCCAGAAGTTGTGTTTGGTTCTTTCTTAATATAGCTATGTTGTCTTTCAAATCTTGGATCAT
TTTTCTGGCTTCTTTGTGTTGGATTTCAACTTTCTCTTGGATCTCATTGAGTTTTCTTTGCCATCTAGATTCTGAATTCT
ATATCTGTCATTTTCAGACATTTCACTTCTGGTTAGGGTTCATTGCTTGGGAGCTAGTGAGATTCTCTGGAGGTGGTAAAA
CACTCTGACATTTTGTATTTGCCAGAGTTCTTGTGCTGGTTCCCTTCTCATCTGAGAGAGCTGATGCTTTTTTTCTTTCTT
TTTTGAATTTGCTATTGTTTGGATGGAGCTTGTGATTTTTAAATTCTTTTTTTCCCTTGTGGGTATGACTGTGGTGTA
ATGTGTATGGGTGGATCAGCTTCATTTCTGAGTGCTTTCAGGGCGCCAAGGCTCTGTATGGGTTCCTTGGTTGCAGATA
AGTTTGTGCGGTGGCTGAGACGTTGCTTCTTGTAGTGATGTAATTTTGTTTTGTAGTGTAATTCAGGCTGCAGCCCAGT
AGGTGGCACTTAAGAGTGAGCGCCAGAAGGTAGGGGCAAGGGCAGAAGCAATGGAAAAGTCTGCAAAGTGCCCTCCTTC
AGCGCGTTTGCCTTCAGTGGGAGTGGAATTGCTGGAGAAGCCCCAAAAGTGGTCTCTTTCAGCCCACTCTCTCGGT
CCGCTGGGGAGAGCCACCTTCGAGTCCGCAACAGTACACTGAAGAGGGATGAGAGAGGTGAGCGATGACCCCTTCTCGA
CAACCATTCCCTGGCTTTGGTGGCGCCCCCTTCAGTGGCTAGCGCAGTGCTCCTGTTTCTTTGACCCAACCTTTGACC
CAAGAGGGGCTTTAGCCGGCTACATCCCCCTCCCTTAGGGGCCGACTGAACCGAGGGCTAGATTTCCTAGGGGAGTGGGT
CCCTTCCCTTACCACTTTTCAGAGCTGGTGGGTACTGTCCCCCACTGAACAAGGAAGCCGACTGGGGAACCCAGTAG
TTGATATGTAAGGGATGCTGGGAAGTGCTGGGTAAAGAAGGGTGAGTCCCTGGCGAGGGCTCCACCCCCAGGCCTGTGC
CCACGGACCTAGGTGAGGACAGGCACTCCAGCCTTTGGGGCCAAATGTTGCATTTCCCAAGACCAACCTAGCCTACCAT
GCCCCCTACCATCCTGTGCCTATAAAAAACCCCAAGACCCTAGCGGGTAGAGACACAAGCAGCTGAAAGTGAGAGGACA
TCGAGGGGAGCACGCTGGTGGAAGAGCACACCAACAGATGCTGCCACCTGGCAGGCCGTCCAGCAGAGGAATGACGCGG
AGTTTGGTCGTGGCAGTCATAGGAGAGCCCGGGCTGCTGAGCGGCTGGACTCCAGGGGGAAACCATCTCCCTTCTGGCT
CCCCCATCTGCTGATAGCTACTTCCACTCAGTAAAACCTTGTACTCATTTCTCCAAGCCCAGGTGTGATTCAATTCTTCC
CTCACGCCAAGGCAAGAACCCGGGATACAGAAAGCCCTCGTCTTGTGATAAGGTGGAGGGTCTAATTGAGCTGGTTAA
CACAAGCTGCCTATAGACGGCAAACTGAAAGAGCCCATGGTAGCACATGCCCACTGGGGCTTTGGGAGCTGTAAACAT
CCACCCCTAGATGCTGCCGTGGGATCGACCCCCACAACCTGCATGCTCCCTAGAGGTACGAGCAGCAGGGCACTGAAG
AAGCGAGCCACTTCTCCAGTTGCACACCTTGCAAGGGGGACAAGGGAACCTTTCTCATTTCTGATGCAACACACGCAGAC
TGTTTCCAGGTCTCAAAGCTGCCCCCTAGCTGCATGTCTTGCCACCCAGAAGAAACCTGGCTTCGGGAACCTCTCCTGCC
CTGCAGTCTTCTAATGCTTTATGTCTCTTCTCTATCGCATCCAAGCATTTCTCTCCTAGACTATCTGCTCAGAAGGTGTC
CACTTACTATTCTGGCGTTTCTCTGTTGGGGAGCCACACACTACCTGCTTCTGCTCAGCCATCTTGATCCCTTTTCATGT
CTCTATTTATGAAGGATGAAAATGGAATGGAGACATGGAGATTGAGGATTGATGATGAGGCTTGTGAAATTTGATGAT
AATTTCACTAAGAACTTAGAAACAGGGCCAGGGGGTAATAAATGAGAGACACAAAGAGAATGGAGTAAGTAGATATGA
TGGACTTGCAAAGAAAAAACCTTGTGTAAGAGCAGACGTGAGCTAATGGACTAAGAGAAGCATGCAAGTGGGGAGACTA
CTTCTGATAGTAAACATTTTGGGAAGAGTGTGCCATCAGGAGAAAGCCCGACTTCAGTGTGGTCAAGGAGCTGAGCAAGT
TATATCCAGAAGTGATCAAGGATATTTGTGAGTTTGCTTACAATATCATAAAGGCAGGACAAGACTCAAGGAAGATTCC
TAGGGAGGAACTGGTGGGGAAGTCAAGGCCAACTGGGCTGGATAAGGTAGAAAGAGCATATTGATGAAAATAACAGC
CTTAGGTGGGAATTGTAGGGGGTAACCTGAAGGGGATTTGAATACAGGAGCCTGTGTTAAAACTAGCTTTTTCTTTTC
TTGCTTTTTTTCTTAGAAATGTATAGAACACATCCATTTATTCCAACACTCCAAGTGAGGAATAAGGTTTGGGCTCAGC
TTGTAGTCCGCACCAATCACTGAGTATTATTTTGGAGGGGCATGTTTACGGGGGAAGGTACGTTGCTGGTGGGATTGTTTC
AAAGTCTTAATGCAAGTGCCCTCAGGTCTATTGCTAGACATTTAGAATTTTCTTGGTTTTTTTGAAGAGAAAGATTAGGG
ATAGTTAATAGTAGTTTAAAAGACTGTGCAACCTTAGTAACATTTCTTTGCAATAATTAATTAATAACACAAGGAAAT
TTTTGCAAAAGTTTGGGCATATAAAATAAATGAAAGTTATGACTAACAGTTCAACTTTCTCCTGATGTCTCACTTTTGT
GAGAACTGATGTAGACCAGAAAATAAATTTCTAGGTGTTTTTTGTTTTGTTTTGTTTTGTTTTGTTTTGTTTTGTTTTG
GGTTGACAGATCTAATTTTTCATGTCAAATTAGGACACTGACTTTCCCTAACAGATGTTTCAGGAGCTGAATTTGTATCCT
AATTCCTAGTCTTGCTAGTATTTATTTTGGAGACCTGCAATCCATCATTATACTAACATGGCAATGCCTAGGAAGTTTAA
TGGCTATACGACTTCTAGAAAGTGCCTTTATTACATATCTGACACATTGCAAATTCACAAGTGGAGTACAAACAATTTTC
AGCAGCTTAGATGACATTTTCTTCTGCCAGTGGTCTGAACTCACCAGGTAGCAACACCCAAAGCATTACATTCTGTC
TCCCTGAGTTGCATTGTATTTTTGATATCACAGCATAAACTGTCAAAGGAAGTATTTCTCTAATTTTAATTGGGGTTAC
TAGGGTCAAAGCCAAGTTTCCAAAGCTCATTTTGTTTTTTATCTTATTATCATGCATTTATAGTAAGCCATTTATTTTTT
GAAGCAATCCTTATTATACTTGGAAAAAACTATTTCTGTAATGTGGTCAGATGTGATTAGATCACTGTTTTATGCTCA
GACGTATGTTCCACCTTTTCCCATAATGCTGTGACATTTGGGGTTTGTATAGCTTCATGAATCGAACTTAAACAGTAAT
GCATTTTCACAAATAAATAGAACTACTCTTCTACCTATTATGATGGATCTTAAATATCCTGCCACATCTCTCTCGTT
ATTCATTGAGTTTCTGATCTTAGTAATTTACCTATACATTAAACATTGTAATTATGTATTTAAATCACCAGGACACCT
ATCATTTGGTCTCTAAAACCTGAGTTGCTGGTTGTTAGAGAAAAATGTATTTAACTTGTATTTATGAATATACTGGATG
AGATTCTGGTTATTTAATAGGCAGTTTAACTTGAGAAACAGAGTAGATGGGAAAAATGAGAACTTACTGTGCTTTAACA
ACTTTACATTAATTTGTTGGTTTCATTGTGAGCTAGTTTATTACAGTATATTGAACGTGTTTACCAAATAGTAACTCGAT
TCCAGTGTTTAGAACATGCTTTTTTAAATGACATTTCTTAACCATTTTGCCTTATTGCAAGGCATTTACGTATATACAT
TGAGAAGCTGTATCTGATCATGTTTCTGCAGAGATGTGTTATGAACTGCTTTAGTACACGGATTTTTGAATCTTAACAA

Fig. 9.236

AGGATTTTGGCCACCTGCAAAATAAGAGGTTTTTCATTCTTTTTTAGAATAGTCTTTTTTCTGGGTCCCCAGTGTATATTT
TGCCAAAATGCCTAAAATATTGCAATAAAAAAACTCAGAGTTTGGTACCAATCTGACAATTAATTTTTTTAAGCCTTTA
AAAGTATATAACAATTTTAGCCAATGGACACCTCTCACTGAATTGACCATATAGATCATATTAGTCCATTTTTCACACT
GCTGATAAAGACACCCTTGACACTAGGAAGAAAAAGAGGTTTAAGGGACTCAGTTCCACATGGCTGGGGAGACCTCACA
ATCATGGCAAAAGGCAAGGAGGAGCAAGTCGCGTCTTATATGGATGGTGGCAGGCAAAAAGAGAGAGCTTGTGCAGGGG
AACTCCTCTTTATAAAACCATCAGATCTTGTGAGACTTATTCATCATGAGAACAGCACGAGAAAGACCCACCCCAT
AATTGATTTGCCTCCAGGACACATGGGAATTGTGGGAGCTACAATCCAAGATGAGATTTGGGTGGGGACACAGCCTAA
ACATATAATAGACCTATCAGTTCCCTACACTAATAAACTTCCCTGGTATTATTTCCACAGCACATGTACCCTGGAAAAT
TAGCTCCAGGTGATGAGAAATTCAGTTGAGGGCAGTAAAAAAGAAACCAGGTATTCTCACACTAATTTGAGGGTTTAAAT
CTCACTGGGCTCCTGTTTGTGTTTTTAGTGCAATTGGAGGAAAATTAGATTTTCTGAGTAAATAGAAATGAGCAATGTTCC
TATGTATATACACTTAACTCTTCAAAAAGAACACATAGGTCTGTACTGGCACAAGTAAAATGTTGTAGAAATTGGGTC
AGGTGCCAAGTGTGAGCAACCTCACTCTCCCTAATCAGACAGCATTCTTATATTATTTCTGGATTAATATAAGTAGGT
AATCTTGAATAAATTCTCTAGTGAATGTTCTACATCATTTATTGTGAGTGAATGTACAGATGGGATAGTCCCAGTAAAT
TTTTTTAGAAAACCAAACCATGTTCCAGGATCATGGATACTTGTATTCAAGGTTTTCATGGGCATTCAAGGAAGAAAGAG
GTGATGCAACATGCCGTAAGCCAGATTCTCTAGAAAACAGACTTGGAGGCAGAGATTTAGGCCCAAGAAATTTGTTTGG
GGGTGGGGTTCTCTTAAGATCAACACCTGTGTGCAGTAGATTTCTCTCTCTTTGGATAAAGGCAATTCCAGGAGAGAGA
CTTAGCTGAGAAGACTCCCAGCTGCTGGGGGAATGAATGCCCTCAATCCTGAATGGGTATCTGGGTTCATATTTGAGCA
AAACATCCTAACTTTAGTAAATAAAGCTACAGCTTTTTCTTATTTCTCCTATCTGCTTCTTGCTGGGCAACCTAAAAT
TTATCCATTATCCCTGCTATCTCACCAGGAAGAGTATAGCGATAAAGAGCAATTTCTATATCTTATACATTATATTTAA
CTATAACATTTATTGTTATTAGATAGTTAGATTAAGTAAGAAATAAGGTATTCATGATTCCTTTTGAATCTGTGTGAATA
TACATATAGTTAAATAGTGACATTGAAAGTATGGTGAATTTTTATATCATCTTTCTGGAGAAATACTGAGGAAAATACT
TGAATTTTGCATAACCAAGGCATCAGTTACTATTCACAATGAGACTACAGAGAAAGCTTTCCAAAATGTTTAAATGAAA
TATGCTTTCAATGTTATGTGGTTTTCTGTGATGTAGCTACTTAATTAGCATTTGTATCAATCACCCTGACATCAAATA
TGATAACTTGCTTTACTGGAAAGTCACTGCTCAATCTCATGAATGAAATTGATGAGGTAGAAGGTAATGGTGCATTGCC
AAGAAAACAAGAAAGAAGGGAGCTAAAAATTGAGAGTTACCTACTTTGCAGTATCTGCTTTATATATGGTATCTGATTT
CAGTTTATCTGAACCCTCCCCTGGAAGTAGCTATTATCAACTTTATGTTATTTACTTACTCATTTTTATTCTTTTTTAAA
AGAATAATGAGATGTTTTAAACATAGAGAAAAGTGCACAGGGCAATATAAAATCACATACATAATCATCTCCCAAGAGT
AACTAATGTCATTATAGGTAAAGTTTGCTTCATATCATTTTAAATATAAATATTGCAGATATGCTTGAAATATATGTTA
GGCCATATGTGTTCAAGCTGAGGAAAGGTAGGTTCCAGAGAGAAATAATCTTGGCCAATCTCACTCTGCTAGTAAGTAGG
GCTAAGATTTGATCGTCTATACGTGGATCGTGGAACTCATGTTCTTTCCCTCCACTATGTATACTTCCCTCTGTACTCCC
TGAAGACTTAATCTGGATAGAGATACATTGCCTAGAGTTTCTCTATAGTGGGAGAAATGAGCCTTTTACTTTCTGTGGTA
CCACGTCAGTTGCCAATTGATTCTGGCACATAAAAATGTATTATATCTTTACCTGTTTAAATTATTTTCTGTACCACATC
AAAGGAACAAGATAAATTTGGTTATAATAGTTTATTTTACTTTAATTTACTATTATTAGTCATTGCAGTACACAAATGC
AGATGTCTGAAGGGATGCACAAAGATGACAAATGAGGCATGTAACATATAAAGAAATACTACATTTCTAAGTGGTTTAC
ATTATGTGTATGAACATGAATTACCTGCATGGATTATTTTAAAGTTCCAGAACTGTTTTTATAGTTGGTTTTCTTTCTTTT
CTATAGCTGTTATTTATGTCTCCAGCTTCATCTTCAAAGTGGAGATCATAATATTTATCTCATAAGACTCTTAAAAAAT
GAGATAAAAGCATTTGATGAAAGCATATGAAAGCATTTGACTGAGCAATTAGTATAAAGAAGCTACTTAGTACTTATTT
GCTATGGTCACCTGAAAAGAAAAGCTATTTTGCATGCAGTGGTAGGGTTGAGAAACAGCCAATATATTAAATATTTTAA
AATATTTTTGTTTAAAGTTCAACCACCTACTTTTTTAGAAAATCACTTAGCATATGAAATTTATTTTAAAGAGGATACTTAC
TCCACAGTTGCTTTAGAAGAAAGAAAAGAAAATCAGTTTGTAAAATGAGATAAATTGTAAGGGTATCTCATCTTTGTAG
CCATATGAGAGAGTCCAAAGAGCTAAGCAAATATTTGCGAAATGATATTTTACTATACTGGAAGTAATAAACAATTAA
AATGTTGTTATTAAGGCAGTGCTGATATTATAATTTTACTATCTCTAGAGGAATAACCAGAGATCTTATTAAACACATGC
TTTTTTAAAGTTGAAATCTTAGATAACTATTATCATTACTATATGTGTAAGAAAACAACGGCATGGAGAAAGATGTGCT
CTATAGAAAACCTGGTAGGTTATTTAAATATAAAGAAATAGGGAGAAAACCTGAAAATTGCATGATATTGTCAAGTCTCCC
TTGGGACTCACCCACACAACAGTTCTTTTCATTGCAAGTTTCTTTTTTCTTAAATGAGTCATTTTGAGTTCTTTTTTAATG
TGGGCACAAGGAAACAAATAAGGATTCTTCCATGCAGCTAAGGCACCCCAAATCTTATTAATTCAAATGTTGAGAGCAT
TGAAGCCATAAAGACAATAAGAATACTGAAACACCATGTATCTGCTTTTCAATTTAAATAGCCAAACAATTCTGTATTTTC
CAAGTGGGCTTCTTAAATATACATATTTTAAACCTTGTAAGGTAATTCTCCAGAGAAATATGGAGGTCTAAAAGACAAA
GTAATTCATTTATCACCTTAAATCTCATTTAATTAGGGAATAATTCTGATATATCAAGAAAGAACTAATTACTGTCT
TCTAGATGCTTGTGCTTTTGTAAAGAAACAAAAGAGCTCAAGCAAGTAGAAGCCTCAGGAAGAGAGCAGTATTACAGAGG
GACCACAGCAACAATGTTTTTTTTTTGTTTTTTTTTTTGTGTTTTTTTTTTTGTGAGATGGAGTCTCACTCTGTGCGCCAGGCTAGA
GTGCAGTGGTGTGATCTTGGCTCACTGCAACCTCTGCCTCCAGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCTGAGTA
CTACCACACTCAGCTAATTTTTGTATTTTTTAGTAGAGATGGGGTTTACCATATTGTCCAGGATGGTCTCTATCTCTTG
ACCGCGTGATCCGCCCTCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCGTACGCTGGCCAACAATCCTT
ATTAATATTAAAATTTAAAATTTGGCTTGAAGTGGAGAACACTTCAAAGCTCATTTACCACAGAACTTTGTCAATTTTTC
TCCTTATTTGACTCATCTCAGTTGGAAAGCACAGATGTTTTATGCGTCTGCCTTGCTCCCCTCTCCCTAAAATTTTAC
TTACCAGCTAAACCGCCAATCCTAATACATCCAACAATATTTTACCATGAAAATACATTCAATTTTTGTATTTAGATGC
ATACAAGTCTCCTGAAATTGTGAAATATTCCAAGTGTACTTGGAAAGCTCTCTGAAGATCCCCCCTCAAAAAA
AAAGAAGATTATCTCTAGTGGTTATATTTTTTGAAGGTTTATTGTTCTATTGAATATATATCTAGTTCTTTTTTCATAA
AAATGAATATCCTGGCTTTAAGTAACTGTTTGCTAAGCACAGTGTGCACAGCTATGCACACACTGATATCACCATTGCC
AATTTTATTAAAAGAACCACAATACAGAGCAAAGAGAAAAGAGAAGTGTCCCTATTAATGGTATTTCAAATGCGTATT

Fig. 9.237

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TAAACAGCTTATCTACTCCAACTAATTTCTGATAGATTAAATAGCTGTATTCCAAATCTTACAAGATGCACATCTCTC
CTCTATTTCTTTCTCGGCTTGCTTCAGCTGAATAGCTCAAATTGTTTGGCACTGGTAAAGCATGAAAATGTGAGATAAA
AAAGGAAACGAAGCTACAGCCTGAGTGGCCTACAGGCCTTAGAAATGGCAACACTTTAAAATTCATTATATTTTACATT
GCTTCAACAGTTACACATTTTTTTTTTTTTTTTTTTTTTTTGGCTCTTGATGATGTCAGAAGATTTATCTGGGATGTCTGT
CTCCACGGGTAAATAGTGACCGGGGTGAAAACCTGAGGGAACTAGGGAGCACCTAGACCCCAAGCCTTAGTAGTCAGCTC
TGGAATGCTCCCTGCAGTGTGGGGTGTGAGTGTGGTGTCTCTGTGTCTCTGTGTACACACTGTGTGTCTGATGTGCAGA
GGGAAGGCAGGGAAGAAATACTATGGCAGACAGTCACTTGTGGTAGCTCTCTAAAGGGATCTGGCACTTCCTGCCATT
TCATGGAAAAGAGGAATGAGAAGGAAATAAGCATTGTGTTACATGCGGGATCTCACTGGATGCTTTCAGTGCTAGAAGGG
AGCTATTTCTCCTCCTAACGTCTGTAACCTACCAGATTTCACTAATTCAGAGGTTGCTTTGAAACTCAGAGTCAAATTC
ATAGCCAGGGACTAAGTCTTCATGATGTGCCTTTTTGTTTTAGCCTCCTTTCCGGCATGTTTTTTGTGTTGCTCTCCTT
CCTGATTACTTTTTATTAGTGCTATCTTAGTTTTATGTATTTTTATAAACTGCCTTAAATGCTTTCTGGAACCACTGTGC
ATTTAATCAAATAGACTAATAACTAAGGAAAATGATAGGGCTGATAGCATCGTCTGACCCATGAGGAAGCTGAATT
TAAGTAGCTTGTCCAAGGTCACCTCAGCAAGTCCCTTGTGCTGTGAGTCTGGTTCTTGGCTTTCAGTCAGTATTTTCAAG
TTTTACCTACTTATTTCTTAACTGAACACGAAAAAACCTCAAATGCTTATTTCTTTAGCTATTTCTATTTCTCTG
CAGTTAATGTCTTATCAACTTGAAATTCCTCCCTGCCTTTCACCTCCAGTAGAAGTTTCGCTCACCTTTCCAGGATG
CTTGAAATACCTTCTCATGAAGCATTTCCTTATCCTGAAAAAGCATCATTGTTACATCTCTTCTTAGACGCTCCTTTGCC
TTCTGTTTTCAATCTAGAAATTTGTTTTCTCCCCCATCTCCATACTGACTAGGATTTTGTTTGCTTGTTCAGTTTAGAT
CACAGGTGTTAAAAAGATGATCTTACATATTTAAATGATGGACATATGATTCCTATTATCTTCTACTTCTATGCATTCC
TGATAGACAACTCTTAGGCTGGAACTAAACAAGAAGGCCTATTAACATTTTTTGCAAAATAAAATTTGTATTCTTAAA
GTTAAGTTCTTGATGCAGGGAGAAAACATTTTATGATTTTCATTTTACTTCCCACAATAATACAAGCATGTCAAGTATT
GACCTACAATAAATGCTTAATAAAAACCTAGTATTTTTTCAGTCACCTTAAACAGGGCATTTGTATATTTAATGCATTAAAG
CTCCATAGGCCATGTCTCCTATGTACTTTGTATGCTCAGAAAGGGAAGTAATGTACACTGGTGGTTAAGAACATATTA
CCTGAGTACATCTTTAAACCTCAGCTCCACTCCTTGCATGTTGTATGACATTGGCGAGTCATTTAACAGCTATGGGTCT
ACATTTCTTCACCTGTAAAATGGGGATAATAAATGTCTATTTAATAAATGTATTGTGATAGTTAACTGAAATAAAAC
ATGTATAAAGCATGCAATAGGGCACATGGGAAGTATTTTATAAGTGTTGTATGTTATTATTAACTGTAAAGAATTCAT
ACAAGAAGGCACAGAAAAACATCAATTGCAACTAAGATGAAAAGTTTGTGCTGTTAGAGCAGGAGTAGTTAAGATGGT
ATGTATTTCAAATATCCAAAAGCTTCTTCTATTTCTGGAGAGCAACCATGGGAAGTCTATTGTGGCATTAGATAGTGCC
ACAGAGAAAAGACAACAGGATGGTGAGGACAGGAGTTGGGAGGGTGGAGCCAAGGAATCAAATAATGTGGTTTCCTAGA
TGACATTCTTTTCAGAAGCCCAGAGGTGCCACATCTATTACAAAGTCTTAACCAAACCATATCCATGAACCTATACAA
GATAAATTTTAAACAGAGTGCAGATTGTCTTTCACCTCTACATCAAACCTATAGAGTCCAGAAAAAATATTTTATATCATGA
TAGTGATAATAATTGAAGGAATCAGAAGGAGTTGATCATTCTCTACAAGCTTGTGTCATGAGGAGGGAGGTGAAAAGC
ATTAAAAAAAAGCTTAAACCTTTTCCCAATATTTCAAAGTAGCTTTGATGGTATTTAGATAGCAAACCTCATTTTAAAG
AGAATAGGATTTTCACAAAACGGAAAAATCAAGCAGCTGGCATAGTAAGTTATCTGAATTTGATCTGAATTGACGTGAG
GGAGCTGGGAATGCCCTGGCAATCCTGCATTCTCTCTCTTTGTGTCAGTATTTTGACAGTAGTGAGAATGTCTGAGTGT
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ATATGAATGTTAATTTCTGGGTATATCTTTTTTTTAAAACTTGCAGTGTGGGTATATAAGCAGGGCCAGATGTCTCTG
TGTTAATCATTTCTACTCTTTAATAAATAACGGTGGCGTACAGTGAAAATAAAGCCCAGATGGCTGACACATTGCACAA
ATCACTGGATGCACTGTATATTCATTTCTATAGTTTCTGAATACTAATAGATGATAGAGTCATCATCTTTTTATCTGTT
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CACCAGATATTGCAGATAGTACCACAATATAATTTGCTTTAATCTCTATGACCTTCAGAATCTATTACATCCTAGTCTT
CCCTAGGAGACTGGGGTTTCTTGGAATTATTTTAAAAAGTGATCCAAAACCTGGATCTGATTTTGCCATGGGGATTGCT
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TGAAGGTGCAGTAACTTTGGGTTTGTGCTGCTGCCATTGACTCAGGGTGGGATGAAAGAAGGGGAACGGACATCATTGCAC
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AGACCCTAGTGGAATTTCCCAAGGGATGTTTTACATAGTTGGGATTCTCAGTAGTTGATGATATTGGGAAGCAGAACATA
ACCAGAGGAGCTGGGAGTCAGCCCTGCCAGGTGGGGCCCAAGCAAAAACCATACTCTGCCTTTCCCAAGTCACAATA
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ATGTAACACTCAGCCTTAAAGGTAAAGGGAGCAGCAAAGGTGTCCCTAAAAAGCAACTTGCATCCTGAGCACTGTGTGG
ATAGAGGCCACTGTTGCCAGACATCCAGCCTTAGCAGGTTGCAGCCTTAGCAGGTTGTGCTGTCTTTGTATGTGGTCAT
TTTGGTACCATCCAGAACCCCACTTACTCTCCTGCTCCTCCACCACCACCTCCAGTCCCTGGTTGCAATGAGCCACAC

Fig. 9.238

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AAAACAGAGGAAAGTAAACGGGCATTACCTTAGTATCAGCAATGCTGGGATATAGTTCTGCTGGTAGCTCCTCCTAAAA
ATACACCCCTAAAAAATAAAATTGTAAGAAAAAAGAGTTAAAAAATAAAAGTTTCAGTCTCCTTGCCCTGGGCAGGTC
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GCTTAGGAACTCTCCAGGGAGCATTGTAAAGTATGCAAAAGTCTGGAACTCTAGGCCAAGTGATTTAGAATCTCTGA
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GCAACTCTGAGTAGATGCCTTTGCCTTGTGCACTGAATACTTTTGTCTATGTTAACAATGCTTTTTTGTCTATTTTATG
ACGTGTTTCTTGTGTATCACCCAAAAGCAACAGATATTTTTGAATAACAGAGGTTACTATGAGCATAACAGTTATGCACA
TCATTGAACCTCAAATTTTAAGATACTTATTACTAAAATGTTATACTGTGATTTATTGAAAATTTTATGAAGAATTCAT
TGACAGGAGCAGGTTATATGTTAAGTGCTACTTTTCTAGTTGAATGTGGCTCAGGAGAAATCTAGTTAACTAAGTCAAA

Fig. 9.239

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TAGATAGTTTTTAATACTTATATTTATTTAAATAGTAGACTTCCATAATCCTCAGTTATTTTATGTCTTCCAAAACCAA
AAATAATTTCTTTAATTTTGTTCATAGATTTTAATTGAAATTGGTGAACATATGCACACATACACAAAACCTGTAGATTT
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CTTAGGGATAGGGTAAAAAACTGAGAGCCCCCAGCCCCCTTAAAGATATTTTCTCTAAACCTCTTAAAGTTAACTCTGAT
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CACCTAGAAGTGACACATACCACTTTTGTCTTAAATAAATTGGTCAAAGTATATCCCATGTTTATGTAGGATACCAAGG
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TACTCCTTTCCAAATCTCCTAACTTTTTTGGAGATTTGATTGATGTTAAGCAGAAAAAGAGGGTTTGAGTCTTATTTT
GGTGAAAGAAAAATATTCTTGGTTAACTTTGTTCATATAATATATCTGAACTTAAAAATAAGTACACAGTGGCATATAATGA
ACTTAAAGAAATAAAATACTAGCAGGAATGTAAAAAACCTGAACATAACACTGAATTGCTCTTGTAGTGTATGCCA
CTTAGTTCCATCTACAGTCTAGCCCTTGTAAATAACACACACACCGTGATTTTAGTTAAATAGGAACATGCCCTTTCTG
TTTCATTTAAATTTCTAGTCAACACAAATTTCTTAGAAAAATGAATTAAGAATGAGTACTATTTCCAGAGTACATTCT

Fig. 9.240

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GGTTTGTCTATAAATTCTAAGCTAAGCTTTATATGGGAACTGTCAAAGGCACAGTTGCAATGTTGCTGCCTTTTTTTATT
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TGATCATTTTTTTATAGGAAGGAAAAGTACCTTTGGTGGATTATATGGAGTGGCTTGGTGTGTTGTCTACAGTCTTCTAAAT
GACATAATTAAGTACAGGCTTTAGATATAATTATTAGTGTATATATCATTTTTAAAATTTCCAGGTGAATTGTAAACATT
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CTTTTTGTGCTTCTGCCTTTAGCGTTGGTTTTGCCAGCCAGCACCAGATCAGGCAGCTCTATTAGCTGCTGCTATCAGC
ACGTGGATGTTCTCTCCTCTTCGTAAATGAGATCTCCTCTACTCTCCCAAAAGTCCCATATTCTTTTGGGACTCTTTCT
TTTGTAGCTCTTGCTTGTCAAGAATGGAATGGGGAAATCATTTCTATGGAGGAAGTTTTTCTTTGGCTTTTGGTAAC
TGACAGAGTGAACAGATTTCTTCGCTGGGTATGCACTGTGCCCTTCTCCACCTATTCCAGAGCTGTCACTCAGGAGCAC
TGTTCAAGGCCATCCCTGCCTTCTTGTGTAGAGGGCTTTCTGTGACCAGCAGACAGTCAAAGACATCGTATCTATAC
ACAGATGGCACATTGATGAAAACCTACTGATGTTTGCCACTTATTATTTATTCTTTTTTCTTAAATATTTACAAAAATTA
CTTTTCTCTTTTGTATGCCACATCTGACTATGTGTGTTAGTTTTTGTTTTTTGTTTTGTTTTGTTTTGTTTTGTGAGA
TGGAGTCTCACTCTGTCAACCAGGCTGGAGTGCGGTGGCGTGATCTCAGCTCACTGCAAGGTCCGTCTCCTTGGTTTCAT
GCCATTCTCCTGCCTCAACCTCCCAAGTAGCTGGGACTACAGGCACCCGCTACCACGCCCTGCTAATTTTTTTTGTATTT

Fig. 9.241

TTAGTAGAGACAGAGTTTCACTGTGTTAGCCAGGACGGTCTCGATCTCCTGACCTCGTGATCCACCCACCTCTGCCTCC
CAAAGTGCTGGGATTACAAGCATGAGCCACCGCACCTGGCCGTGTGTTAGTTTTTATATCTATGTTAGTACCGCAAAAA
TGTCTAAGAAAGCAGACCCTTCTACCAACACTACAATGTCTCATATTGCAGGAGGCCTCATAGTTAGGAAGACAACCTGT
TAGCAAAGCCCTTGTTTTCTCTCATGAGCCACCAGTGTCTGGCATATCTCAATGCACTCTTCATGATGCTCTAAGCTCTA
AGTTAGGCCTATGGTTGCATTGAAGCTCTAAGTTAGGCCTATGATTAAAGTCTTCTGGCTCATAATGATAAAAGCCATT
ATTGTAGGCAGTTAGAACCCTGTTGAAGGACAGAGACGTGATGGATTACAGTCTGAGATAATGTAAGTTGTTTAAAAAAG
TGAAAATAAAAGAAAATCAAAACCTTTGCTTTACCTATTCATTTTTAAATAACCAAGGCATACCCCTTTTGCTGTCTTA
AGTTTCAGACAAGGGTGCAACTTCTACAGTCATTCCTTGGTGGCTTTACATGCTTTATTTGCCATGGGCCATTAGGTGC
TATGGGTGTTTTTCATCCCCACTTTGGTGCTTTGTCAAGACATCAAGCATCTCCATGAATATACTTTAATCTTTCCCTTT
TTTGTGTTTTTGTTTTTTTAGGTAAGTACTTTTGGTATTCTTCTTCTTAATGCTGAAAGAAAATGCAGCTGCCTGATGTA
GAACAAATGGCTTTTGAAGCAAAATCCAAAAGATAAAAAATAATGTGAACAGTAAAGAATGACACCATAACCAGATACT
GGTAAGAATTTTAAGTGGCATTCAAAACACCCCTCTCTTTTTTGGAGAGAGGACTAACAGTACAGGAAGATGCGGGGAGGG
TGGGGAGGGATTGTTGCCGTGCAAGTAAAAAGAAAATTACAGACAGTGTTTCATGAAATATTCATTTACACCTGCTTATT
GGTTAGATGAACAGCCTTTGGTTCTGAGCTGGCTCACCATTCCCAGGCTGGAAATTATTCACCTAAATGCAGCTTTTAA
ACATTTTCTTAAATCCTCAATGGAAACAGATGTGCACATGCTGAGGGAACCTTTTTTTTTGATCTTTTTCTAGAAGAGGG
TTAAATTAGAAAAAAAACCAGTATTTTTTAATATTTAATTTTCATGTGTATAGCTATGAAGCTATATACTTAAATGCTTT
GTAACATATGATCATGAATATATGTATAAACTTCACCCAGAAAAATCCATAAAGCTTTTGACAATAAATGTGTATAATC
TGGATAAATGAAATACCAGTGAACCTCAAACATAATTTATTTTCATGTTTTATGTCTTCTCTTTCTCATTTTCCATGTACA
ATTCTGTGTACATATGAAATAACATACAAAATACAATTGTACGTATGCTTCAATGGCATTTTGCCCAAATTCATAGTC
CTTTTTTATATGGCCGTGGATTTCGAGATTTAAACATATATGGTAAATAGTATGTTTCTGTAACTTCAGCATAGCCCT
TTTAGTAACAAGACATTGTTTACTATAAATTGAGATTATCTATCCAATGTGAGTAAATATATTATTAATAACATTA
CATTTAAGATGAACAAATAGCAAAGGTTAAAAGGTTTCAGAGACCATTACCTGTTACTAAATCATTTCTCTCTCCCAT
TACTAGTCTTATCTCATTGAGTTTCTGTCTGTCTAGAAATGTCATCCCTCTGTCTCTGCTTTATTTAGTTTTTGTCTAT
ATTCAACTCAGACCCTATATCATGTACAAAGCTTTCCTCAAGTCTTTGAACACAGACTGGTTTATTCCTGCCCTAAAT
CATTTTTTAAATAGAATCCATGTGTAAATTTTCTGAGGCTGCCATAACAAAGTGCCACAGACTGCAAGCGTAAATAGTAG
GAATTGATTTACTGATAGTTCTGGAGGCTGGAAGTCTGAAGTCAAGAGAGCAGGGTTGGTTTTCTTCTGAGGCCTCTCTC
CTCAGCTTGTAGATGGCTGACTTCTCTCTATTCTTCACGGGGTCTTTCTCTGTATGTGTCTGTGTCTTCTAGCTCCTC
TTATAACAACGTCACATTTGATTAGGGCCACACTTATGACCTCAACTTAATCACCTCCTTTAAACCCTGTCTCCAAAT
ATAGTCATATTCTGAGGTACGAGGACTTCAACGTACAAATTTTGGAGGTATATGTACGAGGACTTCAATGTATGAATTT
TGGGTATAGGATTCACCCCGTAACAGCCCATGTTAGTTTCTCACTTCATTATTCTCTGATTCCTCATGTATATTGGTT
CTATCTTCCCTAACAAGTTTCATCAAGGATAAGGATTGATCATATTTCAAGTTTTGTTTACTTCATTACAGTTCCTAGA
GGGCTGGCATATAATATTAATGCTACATATTTGTTGTTTGAATTACTGGAAAAAGTAGCATAACAGGACCAAATGAG
AGCCTCTTTTTCCAATTATTGCTGACTTTAGCTCCCTTATCTGGACAGAAAAAACAGTAGCAGAGGTTTGAGTCAGGC
AGTGGTCAAATGAACTGTCTCTGCTTTCTTTCTGGGAGGCCCAACAGCATTCTCGTCAGCAGGAGCATTCTGGCGAAAG
GAAATGCTGATCTCTGCAAATGGGCAAAAGTGTAAGAGCATTGAACCCAGCCTCATACCACAGAAAAACAATTTGGGCT
TGTGGAAAAATAATTACCCAGTGAAGAGTCTTCTGAACTAGAGGCAGGTTTTTGGAGGGGGAAATCATAAACACATTTGT
GGTTGCAAGGTAGCAGATTGGGTGAGCTGAACTAAACAAATTTTGGTTTAAAGAAATTGTATTTTAAGTTCTGAAGTC
ATACTTAATTCTTAATAATTCTAATTGTACACACGCTAATTTAAATGGAAGATGTTTACTTCATTAAAATTTCAACACT
AAAATGCTGGTTTCAAGTTTGTGTTTACTTTCTTCAGGTTTTGTTTATATTCTATTTTTTCATAATTTCTAGAGCCAGCAT
TCAAAAGTAATTTCTGTGCTTCAAAATAGGATCATCCTAAATTCAAATTATATGTAAATGTCATTGAATGCAGATATAA
ACATGGTGCCACAATTTTATAGAAAAATCTAAAAAAACAAAACAGAAATAACCATGTTGCTCTCGTACCAAACGT
GTCTTTGGGTGGGTACTAAGTCCACCTGAGTCTGAGATTCTCCATCTGTAAAAAAGGAATCAGAATACCTACTTTGGG
CAGTTGCTGATATTAAATTTAAATAAGCAAAGCAGTTAGCCAGTCATGTCAAATAGTTGAGCCCAATGAATGGTCTT
TTTCTTTATCTCTCTTCAGGTTGAATTTTGCACAGTAGATAATCCATCTATCACAGAGGTGTGCAGATGTGACTGATTC
ATTGATTGAAACATTCATTGCCCCATTTAAACAATGTTTAGGTATTATTATCCACTGTATCCTGGTTTCATGCTGGATG
TTGGGTAAATAATGATGAGAAGAAACAGACAACCTCAACTTTAATGGAGAAATTGACATTAAAAGTCTAAAAATATGTA
TAAATTTTAAAAAATCAGTGTGTGTAGTAAATATACACAGTACCAGGAGATTTTATAACAGGGAGGGTATGTGTAAA
ATGTCAGGAAATATTTCTCCTAGGATGTAACAATGAGGAGAGTTTCAGAGTATGCATGTGTGCACATATGTGTATGTGC
ACATGTATGGATATGTGTTCCAAAAGTAATAGTGGTTAAGGTGGTGTAGTGGTGGTGTAGGTGGTGTAGGTGGTGT
GGAGAAGCAAGAGAATGAAATAAATAAAAAGATGAGTGTCTGGGAACAGCATGTGCAAAGTCCTGATGTAGGAAACAGT
ACTGAATGGACAGTACGGTTTACCAAGGGAGGGCTGGTGAGGCTGAGATGAAGAGAGCTAACAGGAAGCCAGTAACCTG
AGATGAGGCTAGAGAGGAGAGCAGAGGCCAACTGGAGCAAAACCTTATAGACATATTAGGTCTTCTACCCTAAGAACA
TGGGAAGCCACCTAATTTTAAAGAGAAAGCTTAATGGATGTGTGTGATTGAGTGCAGGGACTGCTACCGCTCTTAATCCT
CGAAATATCCTGCTAACCAGGTTGGCTTTAGTGGGTGGAAAAGTGAGGCCCATACATATTCAAATTTGTATTTTGA
GATTTTTTTTTTGTCTCCTCAATGCTTTTGCCAGAATATAGAAATTCAAAGTATGTGTATTTATAGCAAATACGTAGATG
TTTTTACCTTATAGGTGTGTTTTTATCACATTATTTTTGCACATATGATTCTTGCATGACCCGAGTGAAGGGGTGGGT
TGCCCCCTCCACACCTGTGGGTGTTTTCTCATTAGGTGGAACGAGAGACTTGAAAAAGAAAAAGACACAGAGACAAAGTAT
AGAGAAAGAAATAAGGGGACCCAGGGAACAGCGTTCAGCGTATGGAGGATCCCGCCAGCCTCTGAGTTCCTTAGTAT
TTATTGATCATTCGTGGGTGTTTTCTCTGAGAGGGGGATGTGTGAGGGTCACAAGGCAATAGTGGGGAGAGGGTTCAGCAG
ACAAACACGTGAACAAAGGTCTTTGCATCATAGACAAGGTAAAGAATCAAGTGCTGTGCTTTTAGATATGCATACACAT
AAACATCTCAATGCTTTACAAAGCAGTATTGCTGCCCCGATGTCCACCTCCAGCCTTAAGGCGGTTTTTCCCTATCTC

Fig. 9.242

AGTAGATGGAACGTACAATCGGGTTTTATAGTGAGACATTCCATTGCCAGGGAGGGGCAGGAGACAGATGCCTTCCTC
TTGTCTCAACTGCAAGAGGCATGCCTTCCTCTTATACTAATCCTCCTCAGCACAGACCCTTTACGGGTGTCGGGCTGGG
GGACGGTCAGGTCTTTCCCTTCCCACGAGGCCATATTTTCAGACTATCACATGGGGAGAAACCTTGAACAATACCTGGCT
TTCCTAGGCAGAGGTCCCTGCAGCCTTCCGCAGTTTTTGTGTCCCTGGGTACTTGAGATTAGGGAGTGGTGATGACTCT
TAAGGAGCATGCTGCCTTCAAGCATCTGTTTAACAAAGCACATCTTGACACAACCCTTAATCCATTCAACCCTGAGTTTG
ACACAGCACATATTTTCAGAGAGCACGGGGTTGGGGGTAAAGTTCACAGATTAACAGAATCTCAAGGCAGAAGAATTTTTC
TTAGTGCAGAACAAAATGGAGTCTCCTATGTCTACTTCTTTCTACACAGACACAGTGACAATCTGATCTCTCTTGCTTT
TCCCCACACCCGAGATCTTTTAATATTCTAACATACTCTTATGTTCTAAATCAGTGACACTTAGAAAGAAATGTTGACT
TGAACATCCAAGTACAGGTGCTAATTTGACCTAGCAGAAACATTTTTTAAGGAAATCTCTCTGCACTGAGCACTTGCA
TATCTAGCAATGGAAATTCAAACAATAGAATGATTATCATAAATTCCTTTGAATACCTTTGTTAGCAGAGATGAAACC
TTGGCCACCTGGCTTTAAGGGAAAGCTTAATGAGCATGTGATTTCAGTGCGGGGACTGTTACCTCTCTTAATCCTAGAAA
CATCATGGGAACCAGGTTTGCATCAGTAATGAATTTTTATAATGTGAATCTTTGATTTTTCTTTTCGCTGGTAAGAAACCT
TATTTGGAGATGCTATGTATGTGAGTTTAATCTCATTTTTCCATTCTTTCTTGAAGCTTTCTAGGTTAACACCTGAAAC
AGTGGTTACTATAATCAGTAAAGATTTATAACCAAAGTATAAGGAATTGGGGAATTTCCATATGGTGTGTCTCACAAA
CAAACGTCATGTTTCAATATGATATGAATGCATCTCCATCCAAATTATTATTTTAACCTGTATATTTTTCCCTCAGTCAG
CAATTTATGTAACACATATATTTTCAATTCATCCAACCTCCTAATAAGAGACCCCTAAATTAACCTTGAACCTATGATTTAC
ACCTTCTCAATATACAGTCAATAATTGAATTTAATGACTAACCAAGGACATTTTAGTTCATTGCAACTGCTTACAAGATT
ATATGTAAAGGATTCAGGAATAAATAGAGGGGTCTCTCAGTATGAAGTTTAAATAAATACTTAATGATAAAAGAAAAAT
TTGTCTGCAGTTTTTTTTTTTTTACAATTAACTTTTTACAAGTTATTATCCCTTAGGCTCATTCCATTCTGCTCCCTTTG
TTTTGAAACACTGTTATGACATACTACTGTCAGTAATGGAATGTCAGAAAATAGTACATATGAAAGACACAGTTCATTC
TACTGTTAAATATTACATCATTGAAGGGTTTAAATCCTAAGACGTATCTTTGATTTACCAGCCCAGCCCAACTTCCTAT
TTGCTCTCTGCTCCATTTAGTAGATTTTATGCTGGTTGCTGGACTAAACAAGTCAAACACCTGCAAGGGCCCTCCATCT
GTGGCCAGAAAAGTGTGCTGGTGGTATATTTTTTGATGTTTAGGAAGAAATATTGATCTGCTTAACCTAAGATGGTCATA
AGATAATATGGGGTTGTACTCATCTGATTCTCACAAAACCCAGGGTGCTCAGAACTATACTGATGTGTTGGAGATGCT
ACTTAGGAAATTAGAGGACCAGCATGCATGTCCTTTGGAATGATGTATGCCACCCTATCATCTGCTTGGCCAACAACCT
GAGCCAAGACACACCTGGGGTATCTTGGTACTACCTAGAGACCCCTAGCTAATTTTGCCAGGGTGGATTGGTAGAATCCA
AGGAAATAGTTTTTGCAAGTGACACAATTGGTGGATGATATAATAAGATAATGAAGACTAAAATAATTTGAAGAAGGGAA
ATGGAGATAATTTAGGCTAAGTTGTTCTATTTGCTATTCTTAGAAGTGTTCCTTACATTTAGAAGAAGAAACAATTGA
TTATAAAATCACTGCTTTGATGCATTAATTTGATCATTCTAAACAGGTGATGAATATTGTCTTATGTTATTTGCCTCCA
TTACTTAATCTGACTATCATAGAATAGCTAAGAATACTTCTTAAGAATGAGAGTTTGCAACTACCAGTCACATAGGCCA
GTATCTGTTAACAAAATGCTAGTAATTTTGTTTCATTAAATTTTAAACATTAAATCTATTATTGCATTAAGACCTATTAA
AATGGACAATAAGTAAGGGCCAGATATATCATGAGTAGAAGGAGTCCCTTTCTACTGAGAGCCTATGGAAAGGACAACCT
AGATTCAGAAAAATCTGTTCAACTTCCAGGATGGCATACTAGGAGAAATAAATTTGGGGGTCTAAAGAGTAATAAGCAG
GAAAAAAATCTGGTTTTCTTTGAGGCAATGATTAGTTGAAGCCTACACAAATAAACCAATTACAACATTTTGACTGAAC
TGAGAATAATTACACTGGTAGTCAACTCCTGGGGAAAAATTATGAAGTTCAGGCTGTAAGCTGTCTAGGCTTTTATTAA
CTTGCAATTGTTCTTAACCTGATTACAGGCAAAAAGGTCAAAACATTTTGGCCTACCTAGGTAGCCAGATGGTCTAGAGAT
AATTTCTTAATTACCAAATGACTGTGTTTTATAGTGTCCCTTAGGAAACCAAGTTTTTAAACTGTGTCTAAAGAGAACC
TGATTTAATTTTTTTGATGTTTCTTTTATTTTTTGCTTGATTATATATTATAATCATTTTGTAACAAAAAATCCAAACACT
ACAGAAATATAACTTGGATGTTTATAATCTTCGCTTCTCTCTTAGATAACTATTTCTAAGAGTTTTTGTATACATATTT
AGATTTTTTGATACATGTAAATATATATAATCTTTTTTACTAAAATGGTAATTTACAACATGCTTTTAAACAAACTTTTT
TTTTTACCTAAAATTTCTTAGCTATTTATGTGCTAGTACATGTTGCTCTATGAAGTTGTAAACAGAACAAAGCAGCGTGCT
CCCCAAGAGGACTCTTAGATATTTGAAAGTAGCCATAATGGTAAAGTAAATTCATTCTCTGACTTTGCTTGAGGAAACT
AACATCCTCTCCATCCTGCCTTGAGGACAAGTGTGTTCTTCATGGCTGACACATATCTTGGCAGAGCACATGTGCTGC
AACAGCACACAGTATGGGGTAGGAACAGGATTTGAGTGATTTACTAATCAAGTGTTGGCTAGAAGAGAGGAGACTTGTA
CCACCCCGACCCTGCATCATGCCTCTTGCTCTCCTCTTTAGGGTGAATGTAAAGGGGAATCTGCCAGTTGGGGTACAG
ATAAACCTTTTGGTTACCAGACTGTTGGGGTTTAGATGTCAACTCTTTATAGGAATTGTCTAGGAAAGTCAGCTGTCCA
GGCTTGAGAGACCTTAAGGAAGCATGGAAAGCTGCAGCTCCCTTTTTCCAGGAAAGGCAGTCACCCGTCTTCTTGGAAG
CTGTATTTTCAGGGAGGCCTCTCCAGATGGGCTGGGGGACACTGCCAACTGTTAGCATATTGTCCAGACGACCCAGCATG
AGCTATCTAGGTCATGTTTCAGACAGTACCCTCATGGTGTCTCTTGCTTAAATCATTTGTTCTCTGAGTAGCCTTGAATA
GTAACAAATGTGATATCTTGACATCATCTGGTGGTTGGGTGACAACTGTTTAATTTTCTAACACAGTTGGTTTCTGCAA
TGCCTTTTCCCAAGTAATTTAGTATTGTCTTATCCTGCTATGTCTTCCACAAAGGGAAGATGATAAAATCTTTTTAATT
AGCATAATGTTTGTCTACTTTAATTTTACATTAATAAATTTGTTATACTTTTATTAAATACTTTTCACAGTAAAGTTTAA
ATCCATTTTCTCTATGGATTTTTTCAATTTTGTAATATTTCTTTCCCATTTATAATCATATAACATATACATTTAATAAAG
TATATACAATATATATCAAGCATATATGGTATATATACTATAAATTTTGTCTGTTTTTCTACTTTCTTCTACTTATGTCT
ATTTTCTGTATTTTTTTCCATATGGACATCATTTTTTCACATTTAATGTCTTCTTAACATTACAGGGCAGTTCTAATTGTT
AGAAGGCCTCCTTATACTGATGGCTAGTTTGACTCATTATAAAGTCAGTTATTGTTCTTGGTTTTATGTCTTAAACAAGG
CATCCCCCTCCTCGAACCTTAGAGGAATTGATATCTGACTTCACATGCTCTAGGGGATGCCATGTAAATTTTGTAGAAA
TGTTTAGTATTCTGGCATGTGTGTCTCTAACTTTGTGAGATGATTAGAGATCTGTGATTAGTTAACAACCACACAGGAC
TGAATTAGCTTCCTTCCGTAACCTAAAAGAATTGAACAGATTGAGTAAATTAAGTGGATGCCTCATTAGCTCCACAAAGT
TTATTAGAATTAGAGTCAAAATCTAATACAGGGCTTTGGACTACTAATATCTGCTAGCTACCTTTGGGATGATTTTCACT
ATCATAGGATCATCTGACCAAACAAGCTGTCTCAAATTCCTGTTAGAGTGAAACCTTGGCTGGACATTAGCTTCACCTG

Fig. 9.243

Fig. 9.244

TCTTCTTCTAAGCCCTCCTAACTGTTCCAACCTCTGCCTGTTATCCAGTTCCAAAGTCACTTCCACACTTTCAGGTATC
TTTTCGGCACTGCACCACTCTACTGGTACCAGTTTACTCTATTAGTCTGTTTTTCATGCTCCTGATAAAGACATACCTGA
GACTGGGGAGAAAAAGAGGTTTAATAGGCTTACAATTCACAGGGCTGGGGAGGCCTCACAAATCATGGCGGAAGGCTAG
GAGGAGCAAGTCACGTCTTACATGGATGGCGGCAGGCAAAGAGAGAGCTTATGCAGGGAAACTTCTGTTTTTAAAACCA
TCAGATTTTGTAAAGACTCATTTACTATCATGAGAATGCTGCAGGAAAGACCTGTCCCCATAATTCAATCACCTCCTAC
CAGGTTTCTCCACGACATGTGGGAATTGTGGGAGTTACAATCAAGATGAGATTTGGGTGGCGACACAGCCAAACCATA
TCATGTATCCATCAGAACTCTTGGATGACCAGGTGCACTGTCAATGAGCAGTAATATATTTTTTAAAATCTTTATATAA
GAGCAGTAGTTCTCAACAGTGTGCTTAAAATATCTAGTAAATCATGCTGTCAACAGATGTGCTGCCATCTAGGCTTTGC
TGTTTCAGGCACAAGCACAGGCAGAATAGATTTGGTGTAATTCTGAATGGCCCCAGGATTATTAGACTGGTAAATGAAC
ATTGGCTTCAACTTAAAGCCCTCAGCTGCATTAGCCCCTAACAAGAGGATCAGCCTGTCTTTTGAATCTTTGAAGCCAA
ACATTGACTTATTCTTTCTAGTTGTGAAAGTGCTAAATGGCATCTTCTTCCAATAAAAGTTTGTTTTGTCTACATTTGA
AATTTGTTGTTTAAATGTAGTTTCAATGATCTTCTGGATAACTTCCCTATAGCTTTTGATATCAGCACTTGCTGCTTCACTT
TGCATTTTGTGTTGTGGAGATGGCATTTTCTTAAACCTCATGAGACAGTCTCTGCTGGCTTCAACCTTTTCTTCTGC
AGCTTCTCATCTCTATCAATCTTTATAGAATTGAAGAGAGTTAAGGCCTTGTCTGAATTAGGCTTTTCTTAAGGGAA
TGTTGTGGCTGGTTTGATCTTCTATCCAGACTACTCAAACCTTCTCCACTTCAGCAACAAGGCTGTTTCAATTTCTTAT
CATTTGTATTCACTGGAGTAGCACTTTTTTATTTCTTCAAGAATATTACTTTTGCATTCACAACCTTGGCTGTTTGGTAC
AAGAGGCCTAGCTTTCTGTCTGTCTTGACTTTTGACATGCCTTACTAAGCAAGCTTAATCTTTTCTAGCTTTTTTATTTA
AAGTAAGAGATGTGCTGCTCTTCACTCACTTGAACACTTAGAGGTCATTGTATTAATTGACCTAATCTCAATATTGCT
GTGTCTCAGGGATGGGGAGGACTGAGGAGAAAGAGAGAGATAGGGAATGGCCAGTTGGTGGAGCAGTCAGAACACCCAT
TAATTGAGTTTGCCATCTTATATAGGCATGGCTTATGGTACTCCTAAACAATTACAATAGTAACATCAACAATCAGCAA
TTACAGATCACCATAATAGGTATTTAAATAATTTTAAAGTTTGAAATATTGCAAGAATTATCAAAATGTGATACAGAG
ACATGAAGTGAGCACATGCTGTTAGAAAAATGGTACCCATAGACTTGCTCTACACAAGGTTGCCACAAACCATTTTGT
AAAAAAAAAAAAAAAAAGCCATCTCTGCAAAGCACAAACAAAGCACAATAAAATGAGATATGCCTGTTTCAGTCACT
TTGCTATTTAAAGAGAGAAATCTGTAAAGCCATAGTCCCTTGTCTCATGATTCTTACAGTTTGGACTGGAAGAAATCT
TTAGGAGTAAAGACATTAAATGTGATGAATGTTTTTGTGTTGGCCTTTTACCTATTTATTCTCAGACCACATATTGCC
TTTCTCTGCCCTTTATAGTTGGAGGCTACGCTCTGAAGGCTGTGGTCTCAAGACTTCCATGTCTTTTGTAGATAGATG
TAGCACATTGGAGGCACTGGTGGACATTTGGGGAGAGGCAGAAAGAGAGAAGCCAGAGTGTTTTTACTCTCTTACCTGA
GGCAGCACCTCAGGCAGTAGCTGTGTCTCCTCCATGGCTTACCCTTAGCTAGGTGACCCTAATCCTTGCATACATCAG
CCCTCCTTTTTAGCTCTAGTCCAGGGGCGGTAGTTGCTTCTGTGTTTCTAACCCTAAGTTGCCTCACAGTCCCCTA
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ATGAAGGTTTACTGAGAAAGTAAATTTAGGCTAAACCTCAAGTGTGAATATAAATGTATCAATGGAGAGGGGAAGAA
TATTCTAGATCAGAAGTGTCCAATCTTCTGTCTTCCCTGGGCCACAATGGAAGAATTGTCTTGGGCTACACAGAAAATA
CACTAACAAATAGCTGATGAATAAAAAAAGAATCACAAACAAAAATCTCAATGTTTTAAGAAAGTTTACCAGTTTGT
GTTGGGCCACATTCAAGGCCGTCTGGGCCCTCATGGATTGGACAAGGTTTTTCTAGATCATGTGCAATGGGAAGGAAAA
TGAGGCATATTTTAGAACTAAAGATGTAGACAACCTGGAATGTGACATACAAGAGAAAGAAGGTCAAGAATGTTTGAGG
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AGAATTACTTGGGATGCAAGCAGGGAACCATTTAGAAGATTATTACAGACAGCGGTGTTGGTATGGAGCAGAAATGATAG
CAGAGGCAATGGAGGAAAGTGAGCTTATCTGAGAGATAGAGGTAGGAGGTGGAGTCTGCATGTCTTGGTAATTGAGAGA
CTCTGTGAAACAAGATTGAGGAAAGGTTAGGATTATTTTTCAGGATTCAAAACAGAGCAACTGGGTAGCTGAGGTTCTA
ATCACAGTGAGAAGGAATATTGGAGAAGGAAGAGTTCATGGTTGAACTTCAGGTATATGTCTGGTTATCCCCAGGGAGA
GCTGTCCAGGAGTCAGGTGCGCTGTATGTGTCTGGAGCTCAGGAGGCAGATGCAGACCAAAGAAATTGATGGGGAGCTA
TCCATCTACAGATGGTCATAGAAGTTATTTGAGTCAATAAAATCAACTAAAGGGGGTGCAAGATTAGGAGAACAAGAGA
GTCCTGTCTCAAGCCTAAGAACTGCCAATATTTGAAGACAAAGCTGAAATGAAAGACATGAACAGAGACTTAAGAAGAA
AACCAAGAAAGTTAGGTGCTATGGAAGCCAAAGGAAAGGAGTGTTTCAGAAAGGAGGGAGTGCCAATAGCTTCAGAAA
TGTTTATTGGTTTAGTGAAACACAAGTCAGTTGGTGACCCAATGGGAGAAATTTCCATATAGTAGGAGCATAAACCAGA
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CAGAGTGAGGAGGAGGTAAAGAAGTATCTGGATAGGGAAGTAGGAATTGAGAGGGATCTTATTTTGTACAGTGAAAAATA
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TGCAGAAAATGAAAGTGATTTGATTAATAGTTGTTAGAATTTAAAAATGTGTGCACAGTGTTTGGGTAAAGTTGTTTT
TAAAAAACAGCCACCTTGCCCTTTTATTCTTTATGCTGTGAAACCTCTTTAGAGCACTCAGTCACCTTTTGGCCACTAG
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AGTTGTGAACCTTATTGTTATGAATTGAACTTATTTTAGAGAAAAATCTTAAATAACACATTCCATCTTACTTATAGT
TCATAGGTCAGAGAAAGACTGTGGTGCTACAAAACATTAGCCAATATATTATTGCTTTTACGCTAAGTGTAATGTGT
GTAACATGCTATCTCTTTGAAATTTTTTGCCTTAAAAATGCTAATCAGTTGGCACAAGGCGATCATTTACATAGTCAGA
ATAGAGCTTTTGGTTTAGCATTTTATCTTAAATAAAGGCAGAAATGGCATTGCTCTGGATGTCAGTATGGTGCATTATA
ACCCAAGTGGTGGAAAAATAACTGCTAAATGGCAAACACATAGAACTGAATTCTGCTAGTCAGCTTCCATTTGGTAGA
GATATGTGTGCCCTTGGGTAGCTGCAATGTTAGCTATTATTAATAGTTCAAATCTTTGCTTCATAAAAGTTCTGCATAT

Fig. 9.245

AGTGTGTGACAAATTGAAGTGATTCAAGGAAATCATAGTTCTGTGGAGCTTCCTCCCTGTTTTGTAGTGGAGATTGGGA
ATGGGGGTGGACCATAAAGTAGGTGGTTTTTTTTTTTCTGCCACTCTTAACCTAATTACACACCCTGCCATATCCCCAC
CAACATAAGACTTCAGACTGAGAAAACCTACATAATTTAACCAATGTTAGAAATATAGGCATTTTTTAACGTGCTGAAAAC
AGTGTGTGTTCTTTTGGTTCCTAAACTAAAATCATTTGATACATTTCTCTGTCTATGAAGGAAGCCATGAATGCTATGA
GTAAAGATATTTGTGGCAGGTAAGAAGAGGGAGTTAAGTAAGGGAGTTAATATGAAAAATTCATATAAAGGCTCCAAG
GTTAAATAAGGAATACTTTTGCACAATGGGTTCAGGTTTACTGGGTAGAGCATTTTTTAAAGTTTTATAACGACGTTA
GAAGGAATCGATTTGGGAGAAAACTATATCTGCTTAATGTGAGGGAGCACTGTGGAAAATTTCCAGCACAAACTTTTC
CTGGGCCTCAACATTTCTTGGTAATGAACCTAGATTTGGAGTGATCAAATAATTTGCTGTCTCAAGTAGTAAAAGGGGG
ACTATTACTAACTTCACAAGAAGACAGGCATAAACCCAGCAACATGTACAAAGCAGGATGATCTCAGGTAACCTCAGGAA
GATGCTGGAACACTCTGGGTGAAGGGCATTAAATGCTGTTCACAGTGGAACACAGGACTGCTAATGGTATTTTTATGCTT
TTATGCATCAAAAGCATTGTTCTTTGATTTGTAATCTCTCACATATGTGAAGCAGGGCCTTTCAATCTGGATTTGGGGT
CAGAAGACAACATGTGGATGGGAGGCAGAAAATTATAACAGACTCACCTGGGGAGCAGTTTTTTTGTGTTGTTGATATT
CCCTAGCTGCTGGGAATGGTGGCTCACTCCTGTAATCCCAGTGCTTTGGGAGGCCAAGGCGGGAGGATCACTTGAGCCC
AGAAGTTCAAGACTAGCCTAGGCAACTTAGCAAGACCCCATCTCTACAAAATTTAAAAAAATATTAGCTGGGCATGGTG
GCATACACCTGTAGTATCAGCTACTCAGGAGGCTGAAATGGGAGGATCGTGTGAGTCTGAAGTCAAGGTTGCAGCGAT
TGTGCCACTGCACTCTAGCCTGGATGTCCTTCCACCTCCCCCGCCAAAAGGGATATTCCAAAACCTACAACATTCTACT
ACATTATTCTACATTCTTCTTTTTACTCTCCATATTGATACAGAAACAAAACAAAACACATTTGAGACTTACCCTGCT
AAAAGAGTGTTCTTTGAATTGTGGCTGCAATTATTTCTGTTGCAAACTTTAGTGTATTATTGTTATAATGATACTTGAA
GTAATACCCAGATGTATTGAAAATTCAAAAGACTTCATTGCCAGCTCAACCTGCACTCTCAAGCACTGCTTACTAAGGG
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TCTCCTCATCAGTGTTGGTTGAGGGAAATATTGGGGCATGGCATGGAGAGTCCCACCTCACTTGGCAGCCAGAGAATCTTA
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ATGTGAAAGCGATAACTATTTTTTCAATATTCTCTGATGAGACATTTTAGCAATTATGTTAAAATCTGCACAACCAAAAAG
TCTAAACAGAGCGTGAACCTCTGACATTGACTCTGGAAATTACACACATTTTGTTTTACTATTTTAAAAACACACATGA
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TTCTCTAAAAGTGCTTGGTTTTAGAGATGTTTTGGTTAAGAGTTTGACCAGGTGGTAGCAGATTTACTTTTTCTAACT
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GCTAAATAATTAAGGTAGTTAGTAAGGTCTTAGGCTACTTGTAACATAATTTACTCTTAATTTGTCAACCACTTTATGTTT

Fig. 9.247

TCCATTATTGCCTTTACATTCTATGTTACCTAGATTTGGGATTAATTCATCAAATTTAATATGAAAACCTTTAAGTGACT
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CATTCTAAAGCAAAATGCCACTCCATTTAACATTCAAACAGCTTATAAAGAGCTTGGAAATATGAATTGTGTGGGCCTA
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CGCTTAAAGGAATTAGCCAGCCAGAGAAGAAAGAGAAAATGTTCTTGCCAGAGTACCTATCATTTGGGAAGCCCCAG

Fig. 9.248

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TGGGCCAGAAGAACAGACTGTGTACAGGAAAGACACTACCAAGTGTTATTTGTGGCTGATACAGGAGAGGAAGGGAGAG
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ACTTGAGGGTGAAGGTGGGAGGAGGGAGAGAAGCAGAAAAGGTAACCTATTGGGTACTGGGTTTAATATCTGGATGATGA
AATAATCTGTACAACAAACCCCATGGCACAAGTTTACCTATGTAACAAACCTTCACATCTACCCCCAAACCTAAAATA
CAAGATTTAAAAAAAAGGAAAATTATCTACTCTTTCAAACCTTAAATTTCTGGATTTTAACAGTGTCTGTCTGTTTAAAC
CCAAACAGTGTCTGAATTTGGCTACTGAAGAATAAAATGTAGCCCTTTTTTCAGCACACTGTATGTTTACCCAGGTCCCA
GGATGCTTAAATAAACTGGCGTGTCTATTCAAATCCTGGATAAGAAATAATTTTTTCAAATAAAAATTATCTCACAGAA
TACTCTGAACACCTGCTACTCTCATTACCCTGAACACTTGTGGTTTTGTTGCTATAACTCTAGCAAATGGCATAAAGGC
TAGAAAACTGTGGGATAAAGATACAGCATTTCTCTAAGACCCTGCTGCCTTCAGTAGAATTATTTAATATCCTTTCTA
ATTTCTCCAACCTTATTTTTTCACTGTATGAAAAACAGCTTACAAAGAATTAGTAACATTCATATCAATGATTCCATA
GATCTTCGTTCAAAGTGCAGGTAGAAGGTGCATTTCTCAAAGAGTGTTTTAAACGAGGAAAAAAAATGTGTATCATCAT
CAACGTTTTAGTGAATAAAGACATTGCTTACCGTTTTTATGTTCCCTGAGAGGCTAAGTTTCAAGTTCTATCATGAATAGTA
ATTTATGAATAAGAACCACAATTTTTTTTACCAGAGAATTGGAAAACCGCCCATACATTTCCATATAACCCATCTCATT
TTCTAAGTATCTATAACAGTTTAGTGAACAATATTCCTCTTCGAAAATATAGCAAATAATTATTTCTTCTCCTATCAGA
TATGCATGCTTTGTTTACAGGTAAATAACTCTGATTACCAAACCTACTATTACATTAGGTTGATGTTCTTTTCAACGTTA
GACAAAAATGGATAAAACCTTGCTGCCTACTCAGAGATTTGGTCTGAGTGGAATAGGCTTTTGTGGAGCTACAGAATT
TCTGCTTTATCTACTCAGCCAATAATTGGTCAGAGCATGAGCCTGGTTAGAAATAAGCAAAAAGCTTCTTGTATCCATG

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TCATGTTTCATAAAAGTGGACTCAATAATGGGCATGAACCTGCACGAGGGAGGCATTGCGGAGAAGAATACGTCCCATT
TCTGTACCAAAGAAAAACAAGTACACATTGCAAACAATAAATCTTTATCAAATTC AACACCTTATTTTGAACCTCTATA
ATCATTCAAACGTGGCCTAGACTAACATTTGCTTTTTTATAGCTTTTATCAAGAGGAGTGGAGGTATTAAAATTATTAT
TGAGGGGTGTAGTGATTTTCAACTGACGCAATTCTACCCTCCACCTCTCATGCAGGGGACATTTGGCAATGTCTAGGGA
CTTTTTTTTATTGTAATAAGTGGGAAGTAGCTGATGATTTCAAGTAGGTAAAGAATAGGGATGTTAAATCTCCTATAATA
CACACGACAGCCTCTCACAACAAAAAATTATTTGACCCAGAATGTCAGTAGTGCTAAGGTGGAAAAAACCTTATGTGAA
TCAATTAAATCAGGGAGAACTTCAGAGCTTTTTTAAAGACCTTTATTTATATCTAGATGATTGATATTTTTTTAGGTATTC
ATGAAAATACTTTCTTTTACCTTTCTTGTGGTAAGCACAAGATAACACTTTCTTGCCTGGTTAAAAATGGACAACCTGCT
ACACTTTTAAATAATATAAAGCATTTCAGTAATTCAAACCATCCTGTCTTCTGATTTGTCTGAATTAGTGTGGCTTTAC
TGCATTTTCAGGGCTTATTATTCTTTTCAGTAGGGGAGACTACTAAGATTTTATTAAAGATAGCTGAATAAATGATCAAAT
ACATTATTGTAGCTCCAGACTAGGTAATAAACATTGAGATATGCTTTTCAAGTAGTGGTGAAAATACTAGGCAAAATTA
CACATACACTTACATATATTAAGCGACCATCCTGTTGGCCTGGTATGTGAAGCTCTGCTGAACCTTGCCTAAATGCAT
GGACCCATCGATTGTGAATGTGTGACTACTTGTGTGTTTTTCATCATAACCAGCTCATCCTAATAGCAAATGATATGGTT
TGGCTGTGTCCCCACCCACATCTCATCTTTAATTATAGTTCCCTTAATCTCCACATGCTGTGGGAGGGACCTTAGTAGC
AGGTAATTGAATCATGGGGGCAGTTTCCCCATGCTGCTCTCGTGATAGTGAGTGAGTACTCTGAAATCTGATGGTTTTA
TAAGCATCTGGCATTTCCTCTGTTGGCACTCTTTCTCCTTCCTGCTGCCCTGTGAAGAAGATGCCTTTCTTCCCCTTTG
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CCCCATTCTCTTCTCTATGTAGTGCTGCCAACTCAATAGATTTGAAACAAAATTACCCAGGGACCTTCCCCATCTCC
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ATCCAGTTGCAGGTACACTCTTCTCCTTCACCTCACCTTTTGATGCCATCTCCACTGCTACCCTCTTGGGTCCAACCC
TCATGTTATCTTGCCTGAAAACCGCTAACTTTATAACTAGTCTCTGTTCTATCAACATCCTCTTCTGACTGTCATCCA
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CCTCCCTAAGAGTGGCATGATGGAAAGAGTTGTTTCAGAAAAAATTGGCAATGGAATGCTGCATTAATTGGAATGGAA
AAGAATGAATCCCTACACCAACAGAAGGAAGCAGTGTTGAAATCCTGACAGGGACAATTTTATTACAATTACAAAATACA
TAAATATGTGATTACAATTGTGAGAACTGCTCAAAAGGAAACAAGGACCCAGTGAGAGTATAAAAAATAAGTACCTAACA
TAGTCTGAGTGTTTGGAGAATGCTTCCCTGGGGATGATGAGGCCTGAGGATGAGTAACAACATACTCTGCTGAGAACCA
CATGAATGAAGCCCTGATGTGTTAACTAAAAGGATAAGGTGTCTGAAGTGTAATGAGGCAGAGAGGAGAGTGGCAAAA
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GTAATCAGACAGTAAATATATGTGTACCCTTTTGGAGAAACATTAAATTCATTCCATTAGCAGGCAATGAGTTCTTTAT
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TGCTTGGAAATTATTTTCAGTTGTTCTCATGGCCAAAATTGAACTCATCATCTTACTCTTACCCTCTGGTCTTTTTCTTC
TATTTCTTTGTTGGATTGATTACTACAGAGTTATCTAACTAGAAACACAGGAGTCACCTTAGTCATCTTCTTTTATCT
TAACACTGTCTCTCATCATCAAGTCTTCTTTTACCTATAATGTTCTTATATGAGTCCCTTCCACTCTGTCTTTATTAAT
ATTGCCAACTGATGTGAACTACTGTAAGAGCCTCTGAAGTGATCTCCACATTGCTGGTTTTTCATATAATCCACCCAAA
GGTCTTTGTGTTTGTGGTTGGTTTTTCATTGCTATAGCAATATTCACATTTCTGCAGTACTAGTAAATTACACTAAAC
GTACTGTGTTTTTATAATTGACCTCGAGTATAAACATACAAATTCTACTGCTCTGATGTTTTCTTTCTAAGCAATGGTTGA
ATCAAAAAACAATGAATTTGCCTAATTTTCATGATGAAAACGTTCAAATTTCTCCTAAGATTTCTTGAAATCCAAGCTT
GTGATTGTATGAGAATTCACAATAAACAGCTCACAAAATGTATAAACTTCAGTTTGTGCATAGTCTATGAGGAATTACT
GAAGCATACGGCATTACTCATTTGATTTTTTAATTCAGGCAAAAGTTAGAAATACACAAACACATACGCGTACGGAGGTC

Fig. 9.251

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TCAAAGTCAGTTGTGCGATAATAAATTATTTAACTCACCTTATGATAGGTAAACAGTTTTCTTAAATCGCCCTCCTGC
CCCCTGCTACCCTGACTGAAAATATGTGCTCTGGTTTGTGCTGTTATTTTTGAACAATGTATGCCGGTATCCGATGTAA
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TTTTGATTTGCAAGAGATTTTGTCTTCTCCTGATGTTTCATTTTCGTGAGCAAGTCTTCCCTCTGGGGAAAAACCCACT
TGAATTCTAAGGCTGATAGATGCTGGGAATCCCATATGATGAGTCCTGTGGAAGCAGGACATTCCAGCCCTGGGGTTGC
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GCAGTTTTTAAAGGTAAATTGCCATTTTACTTACCACACTGGATTCTCCAAGTCAGACTAGGATTTGGGTACAATGG
GGATCATTGGGGTTAAATTACTTAAGGGATAATGAGATTTACACAGACCCATTAATCTTTCTAAGCTATTGAGAAATTT
TACACGTACTGTGACTGAGGAGAACCTGATACTGTAAAAGAGCAATTCAGTACAGTTTCACCATCCAAGGACTTACCGA
TGCAAATTCAAATACACGTGCTAAGTAAAATGGGAGAGATAGAGCAAGGGAGATATAAAAATTCCAAYAGAGCAATTC
AGATGCCCCATCTGCCACCACGTGCAATGGATCTATGTTCACTAGTAAGTGTGATTGAGGTAGGAGATGTGGATCTACC
ACTCTTCCCATCTCAGTTCCTTTGGTGAAGTGTGAGTGTGAACATTTTGCCTTACATTGGGTGATTCAAGGGGTCTC
CACGGTAAAAGTGACTATGTCAGATTCTTGCCACATAATCTAAGAGATGACTCCACTGAAGTTTGTGTTACTCTACCAT
CAAAAGATGTATCTATGTTTACAGACATTCTTTTATTTCAATTTGAAAAGAAATTTCAATGTCAATAATAAATATTAGCC
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CAGCAGATTCAAAGGTCTTACTGAAGACCATAAAGTTAGTTAAATGTGGCCAGGTGCAGTGAGTCAACACCCATAA
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CCTCTACTAAAAATCCAAAAATAAAAAATTAGCCAAGTGTGGTGGCATGCACCTGTAATCCCAGCTACTCTTGTGTCTGA
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AACGTGGAATGGTCCATGCTAGGAATAAAGTAGATAGCGGCGAATGTTTGCTAGAGACATTGTGATGGACTGATCTGCC
CTACTATTACATCCTCAGCAATAAATATGAATGTTTAAAGTGGTACAATTGCCAGAAATCAGCCAAAGTTTGGCATAATT
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ACCATTTTATCTTTAGATACATTATTTTCCCAGGAGCTTGGGTATTTTATTGAAGCTGTTTCAAATGCATTTAATGTCC
TTTGTAATGGATTTCTTTATCTCTTCCCCAATGCTCTTGGCTGGAGATGTTACTTTTTTATTTGCCCTTATCGGTGTGAGC
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GCACTACACCCCTTAGCTTGCAAGTAGATGTGAATTTTCTGCCTTTGACTTTGTTGAATCTGTGATGAATCATATGTTT
ACTCTGATTAACATAAAACATCTGGATGATCTAACTTTGGGGACACATTGCTTCATATGCACTGAATGCCTGAAAATTG
GTAGAATTTTAGATTCTTTTTCTTTTATAAATGATACTACCCGAATTCCTGCAATACCTAGAGAGTTACAAGTGCTTAGC
TCTGACCTTTTATTCCATTCAATTGAAGTTGTCCACCTTTAGTTTATTACACATATGACTCTTAGTAGAGCAACATCT
GGATTATTGTCAACAGTTCTCAAACACACCATTTTGGATTTCACTATCAGACTACGAACGAACCCCATGGAAAAAATT
CAGGCATACAGGCTACACCCAGATCCTGAATAGCCCTGGTTTTCTGGTTACTATTTTCTCAGGCCAGATCCAAGAAGT
CCTCTTTGGGCTTGTCTCTGGGATTCTCTGATAAAATTGGCTTTAGATTGAGACTGACGTGAAGATAGAGCTGGTCATT
GAAAGACAGAAACAGATGTGAATGAAATAATTCTCCTTTGAGACATAAAAAAATGTAAGATATACCAAGAAAGGGGAAT

Fig. 9.252

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TGTCAGTAATAAAGGGTGACGAGAAGAAACAACCTAGAGATGTGAGAGGACCAATTTAGGAAATGGAAAGGAGAGGAAAA
AATCAACAAAGTTGAGAATGCTAATTACAAAGTGCCACAGAGGAGGGCAAAACCAGCAAACCATTCCTTTGTCAAAA
TCTTGAAAGATGGCAATTGAGGAACCAAGAATCTTAAAAAGGGGGGAAGGTGGATGAGAAATCTGCATATGAAGGGGAG
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GGAGATAGAAAAATTGTTTTTCTATAAAAAATTGTCTAAAAAATATCTGCACTCCCATGTTTATTGTAGCACTATTAC
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TTGCCAGGTTTCAGGGGCTCACGCCTGCAATCCTAGCACTTTGGGAGGCTGAGGCAGGCAGATTGCTTGAGGCCAGGAGT
TCAAGACCAGTATGGCCAACATAATGAAACCCCATCTCTACTAAAAATATAAAAAATTAGCCAGTCATGGTGGTGCATGC
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CTCGTGGAGATAGAGAGTAGGATGCCATTTATCAAAGAGTGGGAAGGGTAGTGGGGAGGAAGAATAAGATCTAGTATTT
GATAGCATAAGAGGGTGACTACAGTCAACAATAATTTATTGCATATTTAAAAGCAACTAAAAGAATATAATTGGAATGT
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CTGTATTAAAAATATTTTCATGTGACCCATAAATTCATATACTTACTATGTACTCATAAGAATTAAAATTAAAAATAAAAA
TAACCAAGAAGAATTTCTTCTTCAGAAAAATGGAATTACCACCCAGAGAAAAGACCTATAGTACTGGCATTAGATGGTTG
CCCAATAAAGAAGCCAAGGGTCTTTATTCCTCAATTACCCCATGTATTAGGCCATTATTGCGTTGCTATAAAGAAATAC
CTGAGAGACTGGGTACTTTATAAGGAAAACATGTTTAATTGGTTTCACAGTTCTGCAGGATGTACAAGCATGGTGTCTGGC
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TGAGAGCAGGAGTAAGGGGGTTGTGGGAGTTAGCACACACTTTTAAACAACTAGATCTCAGAGAGCTCATTCACTCACT
CACCAAGCCATAAGGGATCTGCCCACATGACCCAGACACCACGCACCAGGGCCCCTCTCCAACACTGGGGATTACATCC
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GAACACCACCCTATAATGAACCCCAAGACTCTACAAGTCCAGTTAAATCATTCAAGGCTCCAGTCAGTATTTTAGTGT
CTCCCTGTTGAATATGAATAGATAGCCAAGTCTCCACAAAACCTCTGAGGAAATTTCTCAACAAAAGCCTAATGCAAC
AAATGAAAATAATGATGATGATTAGGTTGATATAAAAAGAGTTTAAGAAACAAATAGACGTTTCAGAAAAAATAAACTT
CGTAGCCCTAGACAGATGACAAGTTATCACATTAATGAAACAAAAGGATACTATACAAATTTATCAAGTAAGAACAAAA
GAGAGTTAGAATTTAAAACAATGAAAGAAGATGCTAATAAATAAAATCAGAGATGAAAAGGGAGATGTTGCTGGGCACA
ATGGCTCACACCTGTAATCCCAGCACTTTGGGAGGCCAAGGCAGGCAGATCATGAGGTGAGGAGTTTGAGACCAGCCTG
ACCAATACGGTGAAATCCCATCTCTACTAAAAATACAAGAATTAGCCAGCGTGGTGGCACGTGCCTGTAATCCCAGCTA

Fig. 9.254

[illegible]

Fig. 9.255

CAACATCAGCAGAGCTAAATGTAAATGGAGTTTAAATAGAAAGAGACAAATAACCCACTGAGAACCCCTAACATTTTCAGT
GTAATACCCACAGAGTTCCACACCATAAACTTGGTATTCCAGTATAAAGCTCATCTCTCAATTGCACGCCACCCTCCCC
ATCTTATTCACCTCTCTTGCATTATTAAGGAGTGATCTCCCCATGGGATCAAATAGATAGATAGATAGATAGATA
TAGATAGATAGATAGATAGTCTCTAATATATGTATTAGAAACAGAGTCTGTGTTGCCAGGCTGGAGTGCAGTGGCATG
ACCATAGGTCACTGTAACCTTGAACGTTTGGGCTCCAGCGATCTACAGGTGTGCAGGTCTACAGGTGTGCACCACCATG
TCCAGCTAATTTTAAATTTTGTAGAGACAGGATATCTCTATGTTGCCAGGCTGGTCTTGAACCTCTCTTCTCAA
TGATTCTCCTGCCTTGGCCACCCAAAGGGCTGAGAATACAGGTGTGAGCTACCATGCCAGCCATAAACATGAAATTTA
CTTATGTTTTATGTATACCATGTGCACATAGCCTGAAGGTAATTTTACACAATATTTTAAATAATTTTGTGCATGAAAC
AAAGTTTGTGTACACTGAACCATCAGCAAAGGAGTCACTATCTCATGTGAGTACCAAAAAGTTTGTAGACTTTGGAGCA
TTTTGGATTTTCAAGCATCTTGAATTTTAGGTTTTTGGATGGGGATGCTCAACCTGTATATATGCATACATGCATATTT
ATGTAACATACATAATGGTACCTAAAACCTTAGCAAATACTCATATGTTAGCTATTATTTTTTGTGTTGTTACTATGAAT
AATAACTATTAGCATTTTCTCTCATATCTAAAATGCAGTTGGTAAATGCTAAACTCATAGGAATGTTGTAAAGATTTAT
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TGTGCCATGCTGGTGTGCTGCACCCACTAACTCGTCATCTAGCATTAAGTATACTCCCAATGCTATCCCTCCCCCTC
CCCCACCCACAGCAGTCCCCAGAGTGTGATGTTCCCTTCTGTGTCCATGTGATCTCATTTGTTCAATTTCCACCTA
TGAGTGAGAATATGCGGTGTTTGGTTTTTGTGTTCTTGCAATAGTTTACTGAGAATGATGATTTCCAATTTTATCCATGT
CCCTACAAAGGACATGAACTCATCATTTTTTATGGCCGCATAGTATTCATGGTGCATATGTGCCACATTTTCTTAATC
CAGTCTATCGTTGTTGGACATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATAATGCCGCAATAAACATACATGTGC
ATGTGTCTTTATAGCAGCATGATTTATAGTCCTTTGGGTATATACCCAGTAATGGGATGGCTGGGTCAAATGGTATTTT
TAGTTCTAGATCCCTGAGGAATCGCCACACTGACTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAACAGTGTA
AGTGTTCCTATTTCTCCACATCCTCTCCAGCACCTGTTGTTTCTGACTTTTGAATGATTGCCATTCTACCTGGTGTGA
GATGGTATCTCATTTGTGGTTTTGATTTGCGTTTTCTCTGATGGCCAGTGATGGTGAGCATTTTTTTCATGTGTTTTTGGC
TGCATAAATGTCTTCTTTTGAGAAGTGTCTGTTTCATGTCTTCCGCCACTTTTTGATGGGGTTATTTGTTTTTTCTTG
TAAATTTGTTTGAGTTTCAATTGTAGATTCTGGATATTAGCCATTTGTGAGATGAGTAGGTTGCGAAAATTTCTCCCAT
TTCTAGGTTGCCTGTTCACTCTGATGGTAGTTTCTTTTGTGCTGTCAGAAAGCTCTTTAGTTTAATTAGATCCCATTTATC
AATTTTGGCTTTTGTGTTGCCATTGCTTTTGGTGTTTTAGACGTGAAGTCCTTGCCCATGCCTGTGTCCTGAATGGTAATG
CCTAGGTTTTCTTCTAGGGTTTTTATGGTTTTAGGTCTAACATGTAAGTCTTTAATCCATCTTGAATTAATTTTTGTAT
AAGGTGTAAGGAAGGGATCCAGTTTCAGCTTTCTAAATATGGCTAGCCAGTTTTCCAGAACCGTTTATTAAATAGGGA
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TCTGTTCTGTTCCATTGATCTATATCTCTGTTTTTGGTACCAGTACCATGCTGTTTTTGGTTACTGTAGCCTTGTAGTATA
GTTTGAAGTCAGGTAGCGTGATGCCTCCAGCTTTGTTCTTTTGGCTTAGGTTTACTTGGTGATGCAGGCTCTTTTTTG
GTTCCATATGAACCTTTAAAGTAGTTTTTTCCAATTCTGTGAAGAAAGTCATGGGTAGCTTGATGGGGATGGCATTGAAT
CTTTAAATTACCTTGGGCAATATGGCCATTTTCACGATATTGATTCTTCTTACCCATGAGCATGGAATGTTCTTCCATT
TGTTTGTATCCTCTTTTATTTTCAATTGAGCAGTGGTTTGCAGTTCTCCTTGAAGAAGTCCTTCATGTGCTTGTAAAGTTG
GATTCCTAGGTGTTTTATTTCTCTTTGAAGCAATTGTGAATGGGAGTTCACTCATGATTTGGCTCTCTGTTTGTCTGTTG
TTGGTGTATAAGAATGCTTGTGATTTTTGTACATTGATTTTGTATCCTGAGACTTTGCTGAAGTTGCTTATCAGCTTAA
GGAGATTTTGGGCTGAGACAATGGGGTTTTCTAGATATACAATCATGTAGTCTGCAAACAGGGACAATTTGACTTCTC
TTTTCTAATTGAATACCTTTATTTCTCTCTGCTTAATTGCCCTGGCCAGAACTTCCAACACTATGTTAAATAGG
AGTGGTGAGAGAGAGCATCCCTGTCTTGTGCCAGTTTTCAAAGGGAATGCTTCCAGTTTTTGGCCATTTCAGTATGATAT
TGGCTGTGGGTTTGTATAGATAGCTCTTATTATTTTGAATATGTCCCATCAATACCTAATTTATTGAGAGTTTTTAG
CATGTAGGGTTGTTGAATTTTGTCAAAGGCCTTTTCTGCATCTATTGAGATAATCATGTGGTTTTTGTCTTTGGCTCTG
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GGTGGATAAGCTTTTTGATGTGCTGCTGGATTTGGTTTGGCAGTATTTTATTGAGGATTTTTGCATCAATGTTTCATCAA
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GGTAGAATTTGGCTGTGAATCCATCTGGTCTGGACTCTTTTTGTTGGTAAGCTATTGATTATTGCCACAATTTTCAGAT
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GTTGGTGATATCCCCTTTATCATTTTTTATTGATCTATTGATTCTTCTCTCTTTTTTTCTTTATTAGTCTTGCTAGC
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CCTCTACACACTGCTTTGAATGCGTCCAGAGATTCTGGTATGTTGTGTCTTTGTTCTCGTTGGTTTCAAAGAACATCT
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GGTATCCTTGTGACTTTCTGTCTCGTTGATCTGTCTAATGTTGACAGTGGGGTGTAAAGTCTCCCATTTATTAATGTG
TGGAAGTCTAAGTCTCTTTGTAGGTCACTCAGGACTTGCTTTATGAATCTGGGTGCTCCTGCAGGTGCATATATATTTA
GGATAGTTAGCTCTTCTTGTGTAATTGATCCCTTTACCATTATGTAATGGCCTTCTTTGTCTCTTTTGTGTTGG
TTTAAAGTCTGTTTTATCAGAGACCAGGATTGCAACCCCTGCCTTTTTTTGTTTTTCCATTTGCTTGGTAGATCTTCCTC

Fig. 9.256

CATCCTTTTATTTTGAGCCTATGTGTGTCTCTGCATGTGAGATGGGTTTCCTGAATACAGCACACTGATGGGTGTTGAC
TCTTTATCCAATTTGCCAGTCTGTGTCTTTTAATTGGAGCATTTAGTCCATTTACATTTAAAGTTAATAGTGTATGTG
TGAATTTGATCCTGTCATTTTGATGTTAGCTGGTTATTTTGCTCGTTAGTTGATGCAGTTTCTTCCTAGTCTCGATGGT
CTTTACATTTTGCCATGATTTTGACGCGGCTGGTACCGGTTGTTCCCTTTCCATGTTTAGCGCTTCCTTCAGGAGCTCTT
TTAGGGCAGGCCTGGTGGTGACAAAATCTCTCAGCATTGCTTGTCTGTAAAGGATTTTATTTCTCCTTCACCTTATGAA
GCTTAGTTTGGCTGGATATGAAATTCTGGGTTGAGAATTCTTTTCTTTAAGAATGTTGAATATTGGCCCCCACTCTCTT
CTGGCTTGTAGGGTTTCTGCCGAGAGATCTGCTGTTAGTCTGATGGGCTTCCCTTTGAGGGTAACCCGACCTTTCTCTC
TGGCTGCCCTTAACATTTTTCCTTCATTTCACTTTGGTGAATCTGACAATTATGTGTCTTGGAGTTGCTCTTCTCGA
GGAGTATCTTTGTGGCGTTCTGTGTATTTCCCTGAATCTGAACGTTGGCCTGCCTTGTAGATTGGGGAAATTCTCCTGG
ATAATATCCTGCAGAGTGTTTTCCAACCTGGTTCCATTCTCCCCATCACTTTCAGGTACACCAGTCAGACGTAGACTTG
GTCTATTACATAGTCCCATATTTCTTGGAGGCTTTGCTCGTTTCTTTTATTCTTTTCTCTAAACTTTCCCTTCTCA
CTTCATTTCACTTCATTTCACTTCCATTGCTGATACCTTTCTTCCAGTTGATCGCATCAGCTCCTGAGGCTTCTGCAT
TCTTCAGGTAGTTCTTGAGCCTTGGTTTTTCAGCTCCATCAGCTCCTTTAAGCACTTCTCTGTATTGGTTATTCTAGTTA
TACATTCTTCTAAATTTTTTTTAAAGTTTTCAACTTCTTTGCCTTTGGTTTGAATGTCCTCCCGTAGCTCAGAGTAATT
TGATCGTCTGAAGCCTTCTTCTCTCAGCTCGTCAAAGTCATTCTCCATCCAGCTTTGTTCTGTTGCTGGTGAGGAGCTG
CGTTCCTTTAGAGGAGGAGAGGCGCTCTGATTTTATAGAGTTTCCAGTTTTTCTGTTCTGTTTTTCCCCATCTTTGTGG
TTTTATCTACTTTTGGTCTTTGATGATGGTGATGTACAGATGGGTTTTTGGTGTGGATGTCCTTTCTGTTTGTAGTTT
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AGCCTCCCAGTTAGGCTGCTCGGGGGTCAGGGGTCAGAGACCCACTTGAGGAGGCAGTCTGCCAGTTCTCAGATCTCCA
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TCCACCCAGTTGGAGCTTCCCGGCTGCTTTGTTTACCTAAGCAAGCCTGGACAATGGTGGGCGCCCTCCCCAGCCTC
GCCGCTGCCTTGACGTTTGATCTCAGACTGCTGTGCTAGCAATCAGCGAGACTCCGTGGGGTAGGACCCTCCCAGCCAG
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GAACCCAGTACCTCAGATGGAAATGCAGAAATCACCTGTCTTCTGCGTCGCTCAGGCTGGGAGCTGTAGACCGGAGCTG
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TGATACTAATGTCAATTGCATGCATTAATGGTAACATTAATTATCTTCTTACACATTGGGCAAAAAAAGAAAATACCAGG
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CCCTTATAGAATGAACTTGTTTTCAAGAGCTCAAAGAAAACCTTTTATCACATTTATCTTCCACTTTTAAATTAAGAAAT
TATTTCTGAAATGCTAAAACCTTTTGTTTTATTCATGTTTTAGAGGAAAGGAGAGGAAGAGAAACCACTGTTACTTTCT
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ACACATATGTTTGTGTAGTTGTGTGTAATTTTGAGGGGGAGAAGAGGAATTTTGCCCTTAATTCTCACTCCAACCACTT
GACCAATAATTGGGAATCATCTCTTTTATCCCTATTTTTTGCCCTTGTTCTATATGTTATTTTTTTTGGAGATGTTCTTT
ATCATAAAGTGCCTTTACGTAGGCAGTATATAAGCAGTATGCAAAATTGAGTCCCCATCTCTGGCATTGATCTCAAAGT
CCTTTTTTTGTTGTCAAATACCTTTGAATCTGTGTCTTCAAACCTCTGCTTGGGTTCACTCAAATCAGTATATTTCTTTC
CATATGTAGGTCTTTTGATTTTTGTCTAGGTGACTCTTTCCTGAGTTCCCTTTTACATCAGAAATTAACATTTTCCT
AATTATCCAGTCTAGAAATTTTAGTCATCACTTTTCTTTTCACTTCACTTTTAAATATCCAATCCATCCTCAGGTTT
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GAATTAATGGAATATATATATATATATATATAGACACACACATATATATAAAGGGTAGTTTATTAAGTATTAACCTCA
CGTAATCACACGTTCCACAATAGGCCATCTGTAGGCTGAGAAGCAAGGAGAGCCAGTCCAAGTTCCAAAACCTGAAGAA
CTTGGAGTCCAATATTTTCAAGAGAAGGAAGCATCCAGCATGGGAGAAAGATGTAGGCTGGGAGGCTAGGCCAGTCTTTCT
TTTCACTGTCTTCTGCTGCTTATATTCTAGCAGCGCTGGCAGCTAATTAGATTGTGCCCCACCCAGATTAAGAGTGGGT
CTGCACACTGACTCAAATGTTAATCTTCTTTGGCAACACCCTCACAGACACACCCAGGATCAATACTTTGTATCCTTCC
ATCCACTCAAGTTCACACTCAGGATTATCACAGAAGGCATACTATGTTCTGAGCTTTCTTCTCTGTACTGTAAAAAAA

Fig. 9.258

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TGTAAATATCCAGGAAGGAGATACATAGGCAAAATATTCCAACTTTATTAAATAGTTAACTTTTTGTTTGCTTACAGA
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TGCAGATTGCTTGGACAAATTTTCAGCAGGATAGTGAGAAATTGATTTCTAGTGTAGAATATTCTGGGAAATGAGATTTT
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CACCTGAAACAGAGTTAACATGTTTTATTCTTGGTACAGTCATGCTGGTTTTTAAGTGTTTAGCAATTCATTTTACCT
CTCCAGCTAAGGTAAAGGAATTCCTGTAGGAGTAATAGAAGCCCTGTGATTAGTATAAACTTAGATTAAAGATGCTTTG
TGCAGCAATTCCCATGGCATATTTCTTTTTGTCTTTTCTCTTTCTTTCTTTCCCTCCCTCCCTCCCTCCCTCCCTTC
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CAAGTCATATTCTAAAACATGGCTTACTCTGATATTAAGTCCGCTACAGGGGAATGGTCTAATTACTCTGCAAATGC
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TCATTGTTTGGATTTAGCACCCCAAAAATGTCTTGTAATGCAGATTTCTTGGGTCCAGGGAATTGTTTAAGTTTTTGT
TCTAAAACTTTCTTATGAAGCAGATTACATATGGAAAATGACCCCTATGGACTATGAGAATTGGCCTTTAGAGAATGTC
ACTGCCAGCCCTGTTGGTATCTATAAGAACCAATAAGCATTATTTACAGAGAGTGATATACACAGTGATAATTAAGAAA
TAACTATATTTATGATGACAATTTTGCCAAATTTACTAGAAAAAATAAAATATCCTCTGATTTGAGATTTATTGTTCT

Fig. 9.259

Fig. 9.260

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GATGCTAGATATTAGACCTTTGTTGAATAGTTTGCAAAAATTTTCTCCCATTTTATAGGCTCACTCTGTTGACAGCTTC
CTTTCCTGTGCAAGAGCTCTTTAGTTTAATTAGATCTCATTGTCAATTTTTCCTTTGTTGCAATTGGTTTTGGTGTCT
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CTCAGCTCACTGCAACCTCCTCCTCCTGGGTTACGCGCTTCTCCTACCTCAGTCTCCTGAGTAGCTGGGACTACAGGC
GCCCCGCCACCACGCCAGCTGATTTTTTTGTATTTTTTAGTAGAGACAGGGTTTCACTGTGTTAGCCATGATGGTCTCGAT
CTCCTGACCTCGTGATCTGCCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGTGTGAGCCACCACACCTGGCCTGTC
AGGACATCTTCACCTGTTCTGTGTCCAAGATGGTATTGCCTAGGTTGTCTTCTGGGRTTTTTATAGCTTTGGGTTTTA
CATTTAACTATTTAATACATCTTGAGTTAAGTTTTGTATATGGTATAAAGAAGGGGTCCAATCTTCTGCATATAGTTAG
CCAGCTATCTCAGCATCATTATTTGAATAGGGAATCTTTTCTCCAATGCTTGTTTTTTGTAGGTTTGTACAGATCAGA
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CTAGCCAGGACTTCCAAAATGTTGAATAGGAAAGGCGAGAGAGGGCATCCTTGTCTTGTGCCCATTTTCAATGGTAATA
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TCAATACCTAGTTTATTGAGAGTTTTTAAACATGAAGCGATCTTGAATTTTATCAAAGGCATTTTCCACATCTTTTGAGA
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CATCTCAAGGATGAAGCCTACTTGATTGTGGTGGATAAGCTTTTTGATGTGATACTGGATTTGGTTTGCCAGTATTTTA
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GAGAGTATGTATGTCCAGGAATTTATCCATTTCTTCTAGTTTTTCTAGTTTATGTGCATAAAGGTGTTCAATAACTCT
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CATGATGTAATGCCCTTCTTTGTCTTTTTTATCTTTGTTGGTTTTAAAGTCTTTTTTGTGAGAACTAGGATTGCAACCC
CTGCTTTTTTCTGTTTTTCAATTTGCTTGGTAGATTTTCTCCATCCCTTTATTTTGGAGCTATGTGTATCATTCATGT
GAGATGGGTCTCTTGAAGACAGCATACCAGTGAGTCTCGATTCTTTATCCAGCTTGCCACTCTGTGTCTTTTAATTGAG
GCATTTAGTCCATTTACATTTAAGGTTAATATTGTTATGTGTGAATTTGATTCTGTCTCATCATGATGTTAGCTGGTTATT
TTGCAGACTTGTTTATATGGTTGTTTTATAGAGTCATTGGTCTGTACACTTCAGTGTGTTTTTGTAGTGTCTGGTAATG
GTCTTTCCTTTCTATATTTAGTGTCTTTTTCAGGAGTTCTTGTAAGGCATGTCTGGTGGTAACAAATTCCTCAGTATT
TGCTTGTCTGAACAGGATATTATTTCCCTTCACTTATGATGCTTAGTTTGGCCAGATATGAAATTGTTGGTTGGAATT
TCTTTTTAAGAATTGTGAATATTGGCCCTCAATCTTTTCTGGCTTGTAGAGTTTCTGCTGAGAGGTCCACAGTTAGTCT
GATGGGCTTCTTTTGTAGGTGACCTTACCTTTCTAGCTGCCTTTAACATGTTTTCTTTGATTTCAACCTTGGAGAATC
TAATGATTGTGTGTCTTGGGGATGACCTTCTTGTGAAGTACCTTACCAGGAGTTTCTGCATTTCTGAATTTAAATGTT
GGCCTCTCTAACTAGGTTAGGGAAGTTCTCATGGATAACACTCTGAAATATGTTTTCCAAGTTGGCTTCATTCTCCTCA
TGTATTTTCAAGGACACCAATGAGTCGTAGATTCAAGTCTCTTTACATAATCTCATATTTCTCGGTTTTGTTCAATTCCTTT
TCATTCGTTTTTCTCTATTCTTGTCTGACTGTCTTATTTTAGAAAGCCAGTTTTCAAGCTCTGAGATTTCTGAGATTTT
TTCCTCCACTTAGGCTGTTCTGTTATTAGTACTTGTAAATTACATTATGAAATTCATAAATGTTTTCAAGTTCTATCAGG
TTGGTGACATTCTTGTCTATACTGGCTGTTTGTCTGTCTGAGTTCCTGCATTTGTTTTATCATGATTTTTAGCTTCTTCCR
TTGGGTTTCAACATACTCCTGTACTTCAATGATCTTCAATCCAATCCATATTTTGAATTTCTATTTCTGTCTATTTAGCC
ATCTCAGCCTGGTTTAGAAGCTTGCTTTAGAAGGGACRCGGTTGTTTGGAGGAAAAAGGCACCTCTGGCCTTTTGGAGTT
TTCAGGGTTCTTGTGCTGATTCTTTCTCATCTTTGTGGACTTTTCTACCTTTAATGTTTGGAGTTGCTGACATTTGAAT
GTTTTTTTTTCTTTTATCCTATTTGATGACCTTGAGGGTTTGAATTGTGGTATTAGGTGGATTTCAGCTGATTGGCTTC
ATTTCTGGAAGATTTTAGGGGTCCAACACTCAGCTCCCAACTTCTGGACTGTGTGCACTAACTCTGGGGGACTTGTATG
AGGCCCAAACCTTGTCTTCTCAYTCTTCAAGTTTTGGAATCCACTCAGCTAGGGGTGCTGAGATGGGACAGCTGCAGTG

Fig. 9.261

AAGTGCTAGTGGGTGTGGGGGTGCCTGCCTCCTTGCAGATGTTTACCAGAGTGGCAGAGGCAATGCAGCTTACAGAGGT
GGGCAGGAGGCCCTTGCTGGAGCCTGTGTGCACAGTCACACTGGAGGTGGTGTGGCTGGGGGTAGGGTGGTGGCAGGC
ACAGGTCTGAATGCCTTCTCTGTGCCCCACAAGCAGGAGTGATAGCTCAGGGTAGGGGAGGATTCAGTGTCTCTGTAC
AGCCACGTGTTTACTTTTCAAATAATTTGATGAACTGAGTGAAAATATTGAATTTAGACTAGCTGATTTTCAATAACT
ATGCTTAAACTTCAATTTATACAATTTTTTTTCTTCAATTTTAAACTTTGAACCAAATTTGGCTTGTAAACATAAACTAAT
AGATTCAATTAGTTATTTTGAATTATTTTCAAAGATGTGTCCAACATTCTAGAGTTTAAAGTTTATATCCTCAAATTAC
AATTGCTGTAAAGTTTACTGGTTTATTTCTCAATGATCATTTGCTGGTTTATGTATATGAAATATTCTCAAACATAT
CAAGGTACAGATAAAGCTATGAGACACGTAAGCTTCTTCTGTTGTCCATTGGGTAGGTAGAGAAATGGATGACTTGGCT
TATTTAAGCTAATTACTAAAGGAAATAGAGAAGCCTTGAGTCCTTTCTGGAACAAAACAGAATTAAATATTCAATTACAT
GAGGGAAAGCATAATTTAGAACAAAATAATGAAATTTCCCTGGAGTCTGTCTTCTAAGAGGTTTGATGGATTCAATTAA
CATACATTTATTTCACTCCTACTGTGCTGGGCAGTCCCAGGGTTATACAGCCCCAGGAGGTTTCAGAGTCCAGTAGAACA
TGAGCATATACCACTCTTTTTCTTTACAATACTCCTTTAGAGAAAAGAAGACTGCAGTTTGCATTTTGGCCTGGGTAAAG
AAACAAATTAGAGATGTAAAGAACTTTGCTAAGCACTACAGATGCCTTTGACGTGTTTATTCCCGTTGACTCATAACT
GTGTCTCACCAGATAACCTATAGTATATTCATACCSAACTTATCTTTTTCTAATACTTTAGTTATTAGTTTTTAGGTA
ACCTGAAGAAGAAATGAATATGAGCTTTTCATATCTATAGTAGGTTCACTTCCAATCCAGAGAAACCATGGACAGACCTT
TTTTTCAGCTAAACAAAGTGTGGTAGTTTAAACATCGCAGTAAGATCAAAAACCACTTAACTGAGTATGAAAAATATTT
GATGATACAGTAGTCTTCAAACCTAAGTTTAAAAGATGCAAATAGTACATATATGCCAGTACATATGCCAGTGTCTTGAA
TACTGGGAAAAAGGAGGATTTCGAGTGGGGGAAAGTTCACGACATTGTCTCTCTGCCTCTAATCTTTCCTGCTCAAGTCC
TTGCAAGTAGAATCAAATATCAATTGTTCTGAAATTAATCTGCTTTTATTATTTCACTCCCTTGTTTCAGARAACTT
GGTAAGTTTTCACTTCTAAATGGAACAAAGGTCCCACTCCTAACGTATCATTCAAGTTCCTTCACAGTCCAGTCTCAAC
CTTCTTTCTCTTTCTTTTCTTAACCTATCATTTGTTTCAATAACAGTAATATACCCTCCAGGAGTATTCAATCTTTTGGC
TTCCCTGGGCCACATTGGAAGAATTGTCTTGGGCCCATATAAAATACACCAATATGATAGCTGATGAGCAAAAACAAA
ACAAAAAGTCTCATGATGTTTTAAGATAGTTTATGAATTTGTGTTGGGCCACATTCAAAGCCATCCTGGGCCATGTAGC
CCACAAGCCACGGGTGGACAAGCTTACAGGTCTAACTCTGATAATTTCACTACACCTTGGATATTCATACTTCTGGT
GAGCMGAGGCTGCTTACACTTTTTTATAGGCCAATATCAATCATAACGTTTAAAGTGTTTAATAAGTATCAATATTTATCT
CATATGGCTCTCAGAACAGTGAATTAATTCATTAGCAAATATATATTGAACCATTACCACTATCTGGCCCTATTCTAG
TTGTTGGCCAAACAGCAGTGAATCAACATGTCTCTCTCTGAAAGAATTTAATTTTTTGTGGGGAGGATAGTACAAAAT
ATATGAAATACATAAACACATTCAAATGTGTGTCTGATGGTTAGTTCTATGAAAAAAATTAGTGCAGAGTAAAGAAA
TAGTAATACGAGTGTGATTTTAGATTAAAAGTCAAGATAGGTATCCCTGTCTGTAAACTGACATTTAAATCTTAACA
AAATGAGTGAATTATGCATTTTTTCAAAGGAAGAGTGTTCAGAAAAAGAAATAGCAAGTGCAAGCTCCTTAGGAGG
AGGCACATTGTGTTTAAAGGAACAGGAAGAAGTTGAGTATTGTGGGAACTGAGTAAGCAAGAAAGAGAATAGAAAAGGAG
GCCATAAAGAGAGAGAGAGGCTTTTGAAGAACCATGGAAAAGAGTTTGGATTTTATCCCAAATGGAGAGGGAACTAGTA
AAGGGTTTTGAGCAGAGAAATGGCATAACATTTTCAAAGGATCACTCTGGGTGCTCTGTGAAGAATAGAGTAGGAGCAAG
AGTAGAAGCAGGGAATGTTATGACGCCACTGCAATAGCCTTGGAAGGAGACAGTAGTACTTGGACCYGGGTGATG
GCAGTGGAGATGGCGAGAAGTGATCAGATTCTGGCAATGTTTTGAAGCAGAGCTAATTGTGTTTGTATGACGGATTTTT
ACATAGAGTGAGAGAAAAAGAGATGAGTCAAAGATAATTCAAGGGTTTTTGGCCTGAACAAATTGGAAATGTGTGTTTT
TATTTATTGAGCTTGAGGAGTTCAAGGGAGGAGTAGGTTTAGTGGTGGGGTGGTTCTGTTTGTGACATTTTGTGTTTGTG
AGTCTGTTTGTGGAAAATGCTTGGAGTGCTGCCCCATTTCCCTCCACCCACTATAGAGTTACCTGAGGCTATGGCAG
ACTCAGTCAGTTTCCCATCTCAGGCACCTGGGGGCTTTCTGCAGTACAAAGGAGCCTGCTGAGACCGTTTACAGAATTT
TTTAACATTTCCAGTGGCAACACTCCAACATTGAGGGTCCAGAGTTGGTGCTCATATACCCAGGTTCCACATTTCTCA
GTGGGACAGTCTTGAGGCATGTGTATATATGGTCCCACAAAAGGTCCCTGGGGAGATTGGGCCCCATCTGCCCCCTGCTT
ATTAATACACACTGTATTAACTTTTCTTCTTTGCTATCCCACTTCTCCATCACCTAACTTTGATTTCTTAGAATCGCC
TTCTGCTCTTCTCCTCAAATAAACTACGCACTCCAATATTTTGCAGTCTCCTGGAAGGAATTAAACCTAAGACTATTA
AATACCCATATGTCAATCTCCAAGAAGGAGATTGATATACTATTCTGAAAACCAAGGCAAAATGTCAGGGATGGAGATAC
AGTTGGGAGGCGTTAGTCTTTTTTACAAGCCTATTCAAAGCCCATTCACAAAGCACTAGATATGTGTAGATTCTTACA
CAGACTTGTTACCGTCTTGTCTGCCTCATGTCTCTTCCACATGAATACAATGATACTCATCAATGAATGCTCTGA
ACTTTCAAGCAAGGCCACAGTAGAATTTCTAGATAGTAGTAGATTGAGAAGTGATGACAATTTGATTGGAAGCTAAGGG
ATCTTGGGGACTTGTCTTGAATTTGCATTTACATAGAAAGCACATCGTTTTTATGTTTGAYACATATTTATTTGTGGTG
GGTTTGGAAAACCTTGTAGCGGTTATGGAGGGGCACCCCAACCCATCCTTTGGTGTGGCCATTGCTTTTTTAATTATGAT
TTGTGCTAAGCCACACATTCTCATTTTACCTAGCTCAGGGTTCTAAGCATGTTTTTTCATGATGGTTTGGTTAGGAAAGT
GAGTAGCAATGAATGAGCTCCTCAATAAATACATAGCACAGACACTGACAGGCAAAAGTGAAGTTTGTAACTCACTTCCCA
ATCAATTGTTCAACTGAATGTGAGAAGAAATCATTTTTCTCTCCTGTTTTGAGTCAGAATATCACATAAAGCACAACTCT
TCCTGCATACATTGTATCATATTCCACAGTAATTTACTTTTTTGCAAGAAGTATCCAAAATTTGGTTTTTCTGCTGAG
CAGTTATCCAGATAATTGCAAATCAGTGGAAATCATTTACATGAATAAAGATTTTTATTCTAATTAACGTGCTAAATCAA
GCACACCAAATGTCTCTAATTTCTTATATTACTCAAATGGCAATATTTTTTGTGAGTACTATCATATGAAATTTCAAGTG
CCTAATGCATTTTCAAATGGCACTCATAAGCATACTAACGTTTTATAAATAAATGTCTCCATATTTTTTAAGATGAATGG
ACAATTTTTGATTTTTTAATRTAGTCCCCAATTTTCAATTTTTCAATTTTCAATTTTACCGAATGGAGAAACAGCATTCTC
AGAGCAAAAGAGCAAGTTTCAAGTCATGGGAACAAACATAGATGCTAATTTCTTATTTTAGGAAGTTCTTTTTTAAAG
TATTATCATTAAGAAGCACAACTCTGTGGAATATTTATGTGCACATTTAATAATAGAGAATTCGGATGTCAAGTCTGTG
TGTTACTCATACCCATACATGCATCCTCAAAAAGCCTTGGAAGTTAATCCTCAGCTGATGAAAGCTAAGCAATTGCTCT
GGAAGGATACTTGAAGTTTTTGGCTCTTGGGGTGAAGGTACAGTCACGAAAACCTCTCTTCAAGAGGGCGCTGTGCA

Fig. 9.262

AACAGCTAGCTCCTCGTGAAGGAAAATTGCCTTGCAGACTGGCACGAAGTGGATTTCTTTACATCTATTAAGTGCTCTGCTCTCTTCATAAAGGAAAATTTCCCTTGCAAACCTGGCACGAAGTGGATTTCTTTACATCTACTAAGAGCTCTGCTTCTTTCTCTCTCTCTTGCTTTTTTCCTTGCCATGCCTCTTCTTAGGCTCTGTAAGGCAGACTTCTTGCTCTTCTCTAGTTATGCTTCTGCAAACCTGCATCATGCTGATTCTATTAATGATTTGATTTTAAATAATACCATCATAGGTGTTAAATGATGTGTATGCCTATCTGGCTTCTCAGGACTTCTCTCTCACTACTTTACATCTCACCAATGTGTAAAATATCTATGAGACATCTTTAATTTAATATGATTCAAAAGTCTTTACTCTTGACCTTTGTTTGAAGTCTAATCAACTAAAAACCTAAAATTCCTGGTGGGAAAAAAGAAACTTGACTACTCTAGTTTTTTTAATAACATGTATTTTCATTTATTCTGATTATAAAAGTCATCTTTATTTATTGTAAGAAATATAAACATTATGGAAAATAAAATTTAAAAACATCTATCATTCTATTACCCAGAAATAACTTGATCAACATTTTGGTGCATATTTTAATGAATAGATTTTTTAATGAGAAATGCTTTTCAGGAGTAATTAGATTCTACTTAGTATTGACAATTTTATAATATGCTTTTAAACAAAGTATTAACATTAACTTTTTTTGGAAAATGCCACTTTCTGCTACTTATAGCTGTATTGCTATTTTTCTCAGAACTTCTGCTGTAATGAGTTTATACTAGAGTTGCAAATACACGTTCTTTGAATGTACTTGAGGAGTCTGATAGTCTGAATCTGAGATTTTCCACAGCTCTTGAGAGCACCTGAGAACACACTAGCTTCTTCCACGATTCTGCAGTACTTGCTCCTTCCCTGATAGGATGCCCTTGTAAGCGTACAACCTCCAAACATGGGCTGAGTGTTTCAGTAATCATTGTGTGTTGAGTTCTTCAGTTTCATGGAGACTAAAATTAGGTTTTCTCTGTCACTCTTGTAATGTCATCTGACCTTGACCTGCATTTCAAATTGCTGAGAAGTTAGCAAGCTGACTTTTACACTAGGCACATCAGAAATAATTGATTAAAAATAGAGTTATTGAAGTTGAATTTTTTTTCATTTATAGAAAAGTACAATGTCTTGTTTCATGACAAACAAGGCCTAACATTTATATAATGTTTCATTTAAATATATCAAAAAGAAAACAAAATAATAAAACCCACAGAAAACCCAAAAACAACCTCACCTATTAAAACATCTCTTGATTTATAAATATTATAGTTTAGTTTTGTACAATATCAGTAACTGTGGAAAGCTCCCAATTATCTCATTCCCAAGTGCTTGACTGACTCATAACTTACCAACAATTTTCTTATTTTCAGAGAATCCCAAAAACATATTACCTAGGACTAACCTGGAATAAAACTTCATGGTTGATAGCTTTTATCAATTAAATGGGTATTGATTTKGATAACTCATAATCTTGAGCATATCTTTAGATTTACTTTTATCCTGAATTTATTTTTTGTTAGTAAGATTTAGAAATTCAGATTTATTTTTGTGCTTTATATGACATAGAAAAAAACAAATTGTCTTTCTGTACAAAGTCCACCCTTCAAATAGACTCCAGAAGAAAACATAGAGACTTATAACTTAGGTTAAACATAGGCACAATCCTAAATAGTATTCAGATGAGTCTTACCTGAATAATTGCTCACCTAATATGATAATCACCTGTTAAGAAGCAGTTTCTTAAATGTCACTGTTGGAATCTAAACAGTAGATACACAGGTGTTCACTGTAAACTTGTTTCAATTTTTCTGTATTTTTTGAAAATGTTTCATAATACCATCTTGGGGGGAAAAGCCTCTTTTTTGACATGGCAAGAACCATATTCTACTGAAACAGATCTTATATGCTCTATCAAGTTTATATTTTAGACATCCCTAATTCAGCTCATTTCTGGACTATTCCAGGACTGATTGCCCTACTAAGGCATGCTTATTTTTTTCTTTATGTGTATAAGAAGGATTCAAAGGGACCGTTTGTCAGTTTTAATGTGTGAGCACTACTGATATGTTTTATTGAGAAAAAGCTTACTGCCACAGATCACAGAGATATTTTTCCGAGGTAAGATTCTTGCTGGTTTGCAAGTTTTGTAAAGGAGGCACCAGGACCCATCTCTGTACCCCCAGCCACGGAGCCCTTATTTAGACCCACACTGGGCTGAAATAGAACTTCTGTACAAATGTATCTCTCTGTGCTCTTTCCACTTTAACAGAATAGAAAGAATTGACTGGTTGAGCCCAAATATTTCCCCCAGGCCAGWGTTACAAACTGGCGTTTCACCAAAAATAACTCATTTGATATGTTTTTAATGTTTGATGATCTGAAAATCATCTTCTTGTTTTTGTGTTGTTGTTTTTAGAGATAGGGTCTTACTCTGTGTCAGCCAGGCTGGAGTGCTGGAGTTCAGTGGCACACTCACAACCTCACTGCAGCCTCGATTCCCAGGTTTCAGGTGATTCTCCACCTCGGCTCCCAAGTAGCTAGGACCCAGGCATGCACCACCATGCCCTGGCTAACTTTTTCTCTTTTTTTGTAGAGATGGGATCTAGCCATGTTTCCCAGGCTGGTCTCCAACCTCCTGGGCTCAAGAGATCTGTCCGCCTTGGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCTTCATACCTGGCTACCTTCTTCTTAATTATATACATTTTTTTCTTAAATAAGAGCTCATGTTATTTTTTATTTCTATAAATATTTTATATTCCTTCTCTTATTGCCCTGATACTTTAAATAAACCTAAGCCCAGTCAAATTTGTATTACCCTTGAGTTTCAGTTTTTGGTTCATTTCAGCCCTAAGTTCTCTTTGGGCATGAATTGTTTTTACCATTGACTACTAGCAGGCAGCTGGCATAGCCTGGGACATGTGAGGTGATGCTAAACATGCATTTGTGAGTAGAAGGAAGGAAGAAAGGAAGGAAGGAAGGAAGGAAGGTAGGTGGGTGGGCAAGAGAGAGGGAGGAAGAAAAAAGGAAGGGAGGCTGGTTGGTTTTGATTAGCTGTTGAACAAAATTCATATAGAAAGTAGCTTACTAGTCTATATCTTTTAGCTCTGTTTTATTTCAAAGCTTTGTGTGTTTTTGTTAGGGTCACTTATAGAATATTTAGAAAAATACTTGCTGCCTCATGCATTTCCACCATCCAGTGCCTGGTGAGACCTAAAAGCCCATAGTTATAATTTTGTCTTTTGACCAAAGTTTGTTTTATTTTCAAATATAAATTCAATCCCTTAAGACTCACCTATAGCATATATACAAACAAAAACACGTCAAATTTAACTCAGAATTTTGCTGGATAAATGATTTTACTGGATATCATCTGACCAATGAAATAACAATTTGTGTGTTCTCAAATTTACTTCTACATTTTTAAGGTAGTAAGTTTAATTAATAATCTCTAATATAAAAATGGCTTATTCTCTAAAATTTAGAGGTAAGCTAAAGTTTCATCAAGGCCCATTATTTAGCACTTTCCAAACCTTGGGAAGGTCACTTAACTCCACTGAACCAGTAGAAATAAAATTAGTGTAACACTGAAAGCAAAAACCAGAACCAAGTCCAATTGTCTTATTAATAATAGCCTAATTGTGTAAATGTCTGTTATAAATTTCTTAGCTGGTTCTGAGAGATAGCTAGTAAATGGCACCCCTAAGTTTTATATATTCTGTTACAAAACAAAAACAATAACAACAATAAGCTGTTTTCTTTTGCTGTTACTCAAACATCTGGATTTCTATTGTAGCCCTAGCACATTGTTACCTTTGGTTGGATGGACTGAAAAGTGTTGTGATTTTTCTCCTATGTAGAATATTCAGAGGGGAAAGTGCAGGGCAGTGCACATGCATACAAACATTGACAGGATCACTTTTCCCTCAAGCTGAACCTATACTCTAGGCCTCCTCTCTCTCTAGTTACCACAGTTGAACCCACTCAGCCCCCTACTTGACAGCCTCAGTCCCACCCAGGCTCACAGGCCAATTTTGTCTTGGAATTAGGAATTGAAAGATCTCAGGTAAGCCAGCCACAATGGAGACAAGATTTAGAGATAAGCAAACTGACTTGGACTCACATGTCTGTCATATATTACCTCTGTGGTGTTAGCCAAAGTTCCTTAACCTCTATGAGTCTCAGATTTCTCATCTATAAAGTGGTGATAATAATGTCTGCTCGTGAGTTGGAAGATTAAGTAACATGTAGATCTTATACCTGACACCAATTGTTTTTCTCTGTGAAAGTTCAAAATTTCTCTTATGATTTAATTCTTCAAGGAAGGTGACCTGACCTTTTTTCATTCCTGTATTTACTTATATTCCAGGCA TGTATTGGTTAAGATATACATTTTTGCAATTAAGAGGGACCTCAATTTTAACCTCAGATCTGCTACTCATTTGGCTGTATGATCATAGGGGCAAGTTACTTCAACTTCGTAAGTCTTCGTTTCCTCAGTGGTGATACAGAATAAAAATAACACAAAGCACACAGATAAATTATTATGAGGATAATATATAGTATACCCTAAAATTCGTGTGCAGAGTCTTCAGAACACTTTCATGTTAGTTACCTTATTCAGTCTCTCAAAAGCTCTTTGCAATGGCTCTCTCTCAAATGAGAACC AAATGGATAAGACATTATAACT

Fig. 9.263

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CAGCAAGGTCAATAAACACCACAGAATAAACACCAAATAAGTGGTTGATATCATCATAGTAAATGAGGAATTAGAAAA
GTTAACCCGACATTATGGTGATGTTTGGTTCACAACTCATTTCAGTGCACAGGAAGTGTCCAAGTAAGGAAGGGGTCTG
AGGATGACAGGTACTCAGGAATGGGAAAGTTTATGACTGTTGGCTCCAACAAATGGATAACCAGATTTAGCAAGATATT
GAACAGTCACGGAAAAATCCACAGCACCAAATTTAGCACATGAGATTTTATTTAGGGATTCTTGGTATTGGGGGTC
TTTTTTCTTTCTCCCAAGGGAGTGATTTTGTCTTCTCAGCCTAGCAATTTTCATCACTGCTTTTGCTTATTTGTAAGCATA
ATTCTAAAAAGTTGTAAGAAGCCATTTCTGTCTTATTTGGTACGTGGCCAACAATAAGGGTAGCATGAGTTGGAGTCA
GTGTTGCAACTGGTTTCCCAATTAAACAACCTGTCTCTAGGATCCCATAAGTTTTCAAAAAATAAGTAGAAACATTTCT
ATATTTAGGAGAAGCTTAGGGTTTTGAGCAATCGAGAAATTCTATCACCTAATTTTTGTAAATTTCTTGCCACCTAAT
TTTTTTTAGATCTTGTAACAAAGTTCTTAGAGGGTTTTAATGCTGCAATAAAACATTAACATTTCTTCATATGATTA
ACCTTCAAGATATAACAGCTAATATTTTATTCTTACACACATGGGATATAAAGATTTCAAATAGAGAAAAGTGTATCT
ATCAAACTGAGGACTGATATTTCTTTTGAAGTATAAARTGGTATGCCTATTTACCATGCTCATGAAGTAACCAATT
GGCAATTTTTAGAACTTAAAGATGTTTCTAACAGTATAAAAAATAGTTTTCTTACCTCTCCCAAAGCTTACAAACATCA
GTAGTGTGTGCTAAATTTTGTCCAGTTTTTCAAGATGTTTTTATATAAATGTCCTTATTCAGCCCTCTCCAAAACCTCT
TTGCAATGGCTCTCAATGAGAACCAAATGGGTAAAGACACTGAACTCAGCAAGGTGAGGCAACAATCTTCAAGGTCAC
ATGGCAAGTTCATGGCATAACTGTGACTCTAATCTCTAATCCCTTATACAGTGTGCTGCCCTGAAAACCTTGCCCAATGGTA
TTTTGATTAAAAAATTAATTAATCAATCAATAGGAAAAAGAACTAAATAAGCCCTACCTAAATTACCTGTTCTGAA
ATTTCTGCCCCCTATTTAGGCACATATAAGAATATAATGTAAATGCTCCATCTTATTTTTAAGATGCCCTGAAAATATTG
GTTTGTCCATTTATTAAACAAACATAATTGTGGTTTTTATGTCTCTAGACTGTTTGCTAAGCACTGAGTACCCAGAGGG
AAAGCAGATGTTATCCATGTTCTCATAGTGTTTAGGTCTATCAGGGGAGCCAGATACCAAATAAGTCATTACTAATGTG
ATGACATTGACAAAGAGGAAGTACAAAGTGCAGTGGAGACCTCACCTGGTATGGGGAGGCAGGGAAGACTTCCGGAGGA
AGTGAATTTCAAGATGTAATCCTATATCCACCCTCCTTCACTCACCAGTCAATAGCTAAGTAGGATTTTTTATTAAAGA
CTGTATTCATGAGTAGAATATTAAATAGAAAGAATTTTTATGACTTTTAAATCTTCATATACAGAGAATTGAACAAGA
CAATCAGTGACACCAGAAGACTTTTACACACTTGTAGAAAATTGTGTCAAATTATTTATTTCAAATGATTAGTCTACA
TTGATTTCAAGATTTTTTAAAGAATTTGAGTCTAGAAATATTTTTTGAATAGTTAGAAAATCTCTTAGGAACTTGTCAG
TCATTTAGTAGATATTCAATAAACATAAAATTTATTTTCTTTTCTTCCATTTCTTCTCCATGATTTCTTCTCTTCAA
TCCCATTTATCCAAGTAGGCTTTATCCAACAAAGCTTTAGAAGGAGCTTTGTTAGTTTAGTGAATAGAAAAGCTTGACT
TAGTTGTTTTTCAAATTGCTAAAAAAATCCTAAATTAATGTAGATGGAAACCTTAAGTGTCTAGTAAAACATTTTGA
TTCACAATCATAGTCATATTGATTTTTTATCATTTAATTAAGCATGCCTCTTATTGATTGTTTTAGTCTATATCCTGA
TTTTAGCTCTTTAATTACAGTTTGTGTCTGTTTGTGTCTGCTGGGTCAATATAGTAATGGATGCAATTGTAGAAGCTTA
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CTTAAATAATTTGTCTCCCATTTTCAGAACATGGAATAATTGACCTTTCTATCAAGTAGACTCATATAAGGTTTGAATG
AATATTTGAAAGCAATTCAAAGAAAATTTTGGTATTATTTTGAACACTTCCATTTTGAGAAATAACTTATAGTTGATT
TTGATAAGTAATGTAATAAAAATCATTTTTTACATTTACTTGTACTGAATGGATATTAGTTTTAAGGAGTACACAAGCC
CCGTTATCAAACCTGCCTTTTGTCACTCTTTTGAATGYATCATTTCTGAAGATAGAATFACAACGACACCCTCATCAG
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TCAAAATACAGCACTATCCTGAAACCCACTATCTGCCTGCTCTGGTTTCTGAGCAATGTCTTTCAACCTAACGCAAC
CCCCTGATGTAAGTCTTATTTCTCTGAAGTTCAGCTCAACCCTCGTCTCTTCCGTATGCCTGTTAACTAGAGACTTG
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TAGTAGCTATCTCTATAAAGGAAATTTCTGAGTTTCTGTGTGACACACCGAAACCCAAATTTGCCATTTGTAACATCTG
GAATCTTGGAACCAAATCCCAGAAATATTCAAGAGGCAAAAAMGATGCCTTATTTAGAGCTCTGCAGAACATAAGAGC
TAAGTTATTGGCAGGAGACGGGCACCTTGTCTGATACTTTAGTGGGGTTAGGACTTACATGGATAACAATTGGATTGGAG
CAATGTTTAGATTTCGAATCTGGTTTTTCCCTAGGACATATAAAATTAGTGTGCACAACCTTGTAACATAATAATTTGT
CTCAGGGATGGAACATCTCTCCTTGCTATTGAATCATGCTTTTACTAAAATTCATCTGGAAAAATAAACGCTTAATAAA
TGACAGTAATTAGAATGTTCAAAAGATGTCTTCTTTGTTCTTTTATACTCTTTGATACCCATGAAGTAAAAATTATAAA
CAGCTTGTGTATTATAAATGCTAGTAAAGATTATCCTAGAAAATCATTTTATTTAATTTCTTATTTTTTCCAAATGT
CTAAATTTCTCATGTCTGTGAAATATTAGCCAGTTCTTCAAAAACAATATTGATATTTCTATTTGAAATCAAGGCTTAAT
TTTGCTCATGATTAATCTCAGAAATAAGTAGCTTATTTCTATGACCTTACTGAGACCTGGTAACCTAAGAAGGATTATG
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CTTGTGTCTGTAAGTGAATTTTACACTAGGAAAATGAAGCAGAGTTTCAATTTTCTTGTTCATAATTTACATTTACAGT
TCCATTGTTTTATCTTGCTTAACTGTTCTAGATGACTGCCTTTACAAAGCAGGGGGAAAAGAAGGAAGAAATGTTTTTG
ACAGTAATGAAAAAGACAAGATTAGTTCTCCTTCCCTTGACATTTTTTAGAAAAAGTTGGACATTCCTTAGCAGATTCT
TCTAGATTATTCGTATGTCTTGTAGCTTATCAATTAGATATCCATTAAATTTGAATATCCCTTTAAAAAGGTAAATTT
GTAAGCAAAGGCAGTTAATTATTTGTGAAAATGTATACTGCTGGCTTTAGCCTGAATACAAAGATAGGGTTTATCTTGC
TATCAGTAAATTTGGGATAACAATGAAAATATTTTTCTGCTTCCAAAGTCGTATACATAAACTGTAGCTTTTATTAGAA
ATCAGTGATGCTGTTTACCTACAAAATTTAAACACCACTGGACAAAATGAGAGCTTTGTGTTGCCTGGAGGGTGAAA
AAAGTCCATCTTTGCCAGCAACATCCCTTCTACTCTTCTGCTTGTCTTTGTGGCCTCCCTCCCCCTCCCTTTTACACCA
AGCCACCTCACCCACACAGCTGTTAGGGACTCCTCCCTCTTTACGCAGATCACCAAGTTATCTTACACTGTCTGGCTTT
AGTCTCAGATGAAGACATTCTAATAAGCACCAAGTTGTTTCAAGTCTAGTGTAACTCTGCCAGGAATGTGTGCATTTCT
AAGAGCTGCCGAAAATGGAAATTAATAGTTTGAAGAATTAATGATAATGGTGGCATTTCATACACCAGAAGGGCAG
ATTGGCATCAGTGTAAGCACCTTTTCAAGTCTTCTTTGAGAACACCAGGAAAGTGGCTACCTGTTTTGCTCATGCGAAA
TTTGGATCTCTATACAAAAGCAAAAAGACAGTTTGTGAAGCGTTACAATATTGCAAGTTGATTTCAGRGAGAATGTGTTG

Fig. 9.264

TGTTTATTTTTGAACTCTCTGCTCCTCCAAGGTTGCTGCTGTTTCAGGGCAGTTCTCACTCACCCACTTTCTTTGAGCTC
ATGTCTGATAAGAAAGAGGTGAACTGTAAAAACCTTTTCTATTACTCGTCTCCAAAGCTGATATGAAACCTGTAGCAT
TCTTAAGAACCCTGGTGTCTGGATGCTGTTGTGAAAACAAGCATAATGTTTAATGTCTTGAGCTTTTATTGAAATTA
TATGAATATTCAAGACTCCCTTGGTGTACAAGAGACAGATTGAGCTTTAGAGGTCTCAAAATTTGCAGATATGGTGATG
TTCAGTGAGCTCAACTCTTGGTCAATTTTGTCCAGTTTCAGAGAGGGTTAAATTTCACTCTTGGGCAGTTGAAGCCTCTCT
AATCTTATCCTGAAGAAGTGGCGCTCTCCCTTGGTTTACAGTTGAGGTACCCGCGGGGCAGTGTTTGGATACAGACTG
ATGAAATTATGCTGCATTGTTAACATTGAATACCCTCAGTGGTGAGGACCGATGACGGCAGTGGGTCCTTTGACTCC
TTGGGTACAATTTCTTGAAATAGATGCTTTCCCAAATGTCCTGGACTCATAAAATATATGAAGGATTCTATTTGGCTT
TCATCATTTTATTTAATTTGAAAGAAAGTTGTTTAAACAGATTTATCAGAGTTAAGAAATGTTTCTAGGGAATAGAAAAT
GGACAAGATAAATTGATCTTTGTTGTGAGAGTCATGCATTGTAATTCAGTCTCTAAGGTGACGCAGCATCTCCTTGAGGG
CAAAGCTGCCAAAAGTTCTGAAAGTGATCTGCCTGTTTTTTTTTCTCACACTGACTTTTAGAGAGGGCGCTGGTATTTTC
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GAAGAGAAAACTAAATGATAAAGAGGAAAGATGGATTGTGAATAGAGTAAATATATATTACTGTAATTATGTTAAGC

Fig. 9.266

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CAATCAAATGCAAAGGACAGTTCAAAGGGGACAAGTATGTAAGCCACATTCCAAAAAAGCTTTGAAGTGA
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Fig. 9.267

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TTGTCTTGGCTCATTAAACAACAATGGACAAGTTACCTTAACTTCATGGGTCCTCAACCTTCTCATTTCGAAATTGAGTG
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GGTTATCAGAACCTAGGAAGGGTAGTGAGTGGCAGGGGTGTGGGAGTGGAGATGGTTAATGGGTACAAAAATATATTTA
GATAGAATGATTAAAGATCTAGTATTTGATAGCACAAAAGACTGATTGCAGTCACAATAATTTATTGTACATTTTAAAT
TAAAGAGTATAATTGGATTGTTTGTAAACACAAAGGATAAATTTATTGAGGTGATGGATACCTCATTTACCCTGATGTGAT
TATTAGGCATTGTATGCCTTATATCAAAATATCTCATGTACCCCATAAATATACCTACTATGTACCCACAAAAGTAAAA
AGTAAATACATTAATTTTACCTGGTTGTAAAAATAACTTTTTTGTAAATATGGCTTTTTTAAAAATTTAAAAATTACATGTTG
TGGCCATCATTCATGGTTTACATTATRTTCTACTGAACAGCGTTGATCTAGACAGTAGACACTACGCAAAAACAGCTA
AGAAAATTATTAATTTCTTTAAAGGATAAAATTTAAAGTGATAAATGAGATCAAATTTTATGAGCAAGTCCATCGTAGT
TCACTCTGTGCTTGTCTTAGGGAAAAAATTTGGAATTTCTGTAGAGTGTGCCAGTCAATGCAAAAGTGTCTCATCAA
GAGAGAATTAACACATTAGCAAAATAGGGGCTGTTGATGAAAATGAGATTCAGAGAGGGTGAGAAGACTGAACACATTC
CTGTAAAGCAACAGGGATTAATGCACCAATGAGAATTGCTTTTTTTTTTTTTTAAACAAAACAAAACAAAACAAAACAAA
ACCCACTAAATTTTTCTAGGGGAAAAGTAAATGCTAGAGTGATATTAGTGAATTAGGGGATTTGTGAAGATGCATTTGA
ATGTCAAGAATATAATGTAGTTTTCTTAGTATTTTTTGAATTCAGTGACCTTTTGTAACTCAAGAGACTGAGGCTAAG
ACACCAACATTTACCATGTGCTTCAATCATCTCTCAAGGACGCACAGCTCCTCTGAGCTGTAATAGGAATTCAGGTCT
GTGTGACTCCTGAGCCCATGACCGAACGTGTGCCCATGGAACACTGGTCCCTGCAACTGCTCTGCAATAATGGTTC
TATACTTAAATCATTTTAAAGAAATGTTGCATGTATCGTTACCGTCTTAAACATGATTGATTGTCATATTAAAGGC
ACTGAGAATTGCTAAAAAATGTTTAACTTTTTTAAATCTTTTTCTCAAATCTTTTTGCCACAGAACACCTCCTCCCC
TGCACATGCATCTATATGCCATTTATCACCAGTTGCATGCTGAACTAACCTTAGACAGAATGCTATTTGGAAATGCTG
AGATGCCTTAAATTGCCTTCTATCATCTCTCATTTATTAGTTACTAAGGAAAATGGCTTTGAGAAAATATAAAATATTTT
AGGAAAATATGTAATTTGATGCTCAGATAACTCACTTTTTTGCTACATCAGAAAAAGCAAAACTAGAAATTTAAAAATAAA
CTTGATGCATGTCTTGGTTATTCTTATATTCATAGTGTGTGATCCAAAGGGAGTCAAAGAAATCTGTAGCGAGGCTGC
TCACAGTCTAGGTCTCTTTTTTGTGTAGGACAAAGGTAGGGCTTGGCTTCTAGTTGAAGCTACAGTTCTGTAGGACTTGG

Fig. 9.268

Fig. 9.269

CTCTTCCAACCTTTTTTCCCCTAATTTCTTCCCTTAGCACTGATATTATCAAACAAGTGGCCAATCCTATAAAGGCTATC
AGACTCCAATTTAAAAAACTATTAAATTTAAAAACACTCAGAGATATAATGATTGTATGCCTTTTATGAGGAACTTA
GTACCTAAAAGAAAACCCCTATAACTTAACATTAAGCATATGAAGGTAGCTATTATTCAAATAAGTAGCAGTAAAGACTT
TTCAGCTTTGGATCCTCATAAACAAGACCTGCAGGTTTAAAGATTTGCATACATCTTTTAAACAATGCATCATTTTTTATT
ATAGTATATTATATACTATTTTATATTTACTTAATGAGATTACTGACCTGGGTTTCAAGATATGTATGAAATATAATTA
GGATATGAAAATTAAAGATAAATTATCCCCACCAGAAATAATCTTAGCAGGTTATTTACTAAAGAATCTTAAATCCAGC
AGATCAAGAAAACTCCCCAAGGAGTCTTTAATTAGAACTCTTTAGGATGGAATCAAAATCTCTCCATAAAAAATGAAT
CTTGCCTTAGGCTACATAAATTACAAAATCTGAAGCCCTTTAAGCAGCATTAAAACTGCTGATTTTAAATTGCTCTGATA
ATGAAGTGGAGAGAGACACACTAAGCTTCCTGGATTTCTGTTGGAAATGAATGGGATTCTTGCCAACTATGGATCACAG
TTATGAGGTATTCTGCACAGGGAAAATCAAAAGAAAGCAGATGGACCTTGACATGCTATGCCATCCTATGCCTTCTTCC
TTCCCAAGAAGCATGTTACAATAACTTTAAGTGAAGGTTGTCACAACTTTAAGTGAAAAGTCTCCCCTTTCTGTCTAAT
TTTTAATGTAACCTCCCCCTTCTCGATATATCCAGAAAATAGTTAATAACAATCCCTGAGGGAAAAGAAAAATGCTGCC
CTTCATATTAAAAAAAAGAAGATAACAGAGATCCCCTTAGTGGTGGTGGATTATATAGTGAAAACAGCCAACCTCATTCTC
TTTTGCCTCCTTTTGACCTAGAGAACTGGATTTCGCGGATTGTAACTGGATTTTGATTATAGAGTATGTTCTATGGCTA
CAAAAGAGTTTTCTTTGAGGAAAACCTCAAACTGCCAACAGATATGCTTGGTATATCAGTTTTCAATGCTTTCAATTGT
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CTGTTGCATGATGTTATCACCATTTGCAAAGAGGCAGCATCTCCACAATTTTCTTGGCCTGTTCTCTTGATGGCAAAT
GGCTACTGCACGTCCAGATGATCATGTCTGTGTCACAGATAGTGAGAGGGAAGGAGGAAGGATGAGAGAGGATGGCCAG
CTAAAGTTGTATCTACCCCACTTATAAGAAACACAAAGCCCTTCCCAGAGCTTACAGCCCTTAGCCAGAACTATGAC
ACATGGTCATCTTGAGTGCAGAAAGATGGAAGTTAACATTTGGTTTTCTGGTCTTTTAAATAAAGATAGCAAGGATGAA
GAAGTTTGGAAGAAAGACTGTGGGCTAGCCAATCAAATTGTCGACCACATTTGGCCTGTATTAATGTATAGTTTTTAAAGC
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TAAAAACAAGATAAATAAATTGTTTGAAATATTTTCATCATATTGCCTCTGAAATGATTGATGTCATACTGCCTGG
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AACTGGGAAACCCACAGAGAAAGGTAAATGAAATAGGAGAGCCAGGTAAATGGGCCATGGATCTAAGAACCAGGAATCA
TGATTGTTTGGAAGAACATATCTCAGACTGAGAAAAGCTGGCGCTAAAGTAAGCTAGCATTTGATTAGAATGTTGAAGG
ATAATTTCAATCTTGGAGACAGAAAACCTAAGTAGAAGCAAAAGAAGTAGGAAATAGAAGCCTGTGTTTGAGAGATTCA
GAGTGACTGTGTTTTATTAGAATTGCAAGAAAGGGAACCTTGAAAACATAAATTTAACATGGGGCAACAGGCCCTTGGATT
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AGTCATATGCTGAAGTCCTAACCTAGGACCTCAGAATGTGGCTTTATTTGAAGATTGAGCCTGTAAATAGGTGATTTTG
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CACACGCAGAGAAGACTACGTGGAGATGAGGAGAAGATGGCCATCTACAAGCCAAGAGGAGAGGTCTCCCGAAGAAACC
AACCCCACTGACATCTTGACCTCAGTCCTCTGGCCTCCGGAAGTGTTTTAAACATAAGTTTCTATTGTTAAAGCCACC
TAAGTCTGTGGTATTTTTGTAATGGCAGCCCTAGGGAAAACCTACAATGAGCAAGTATTATTTGTATTTTAAAGAAATAAT
ATTATAAACTAAATTCATGGAGGGGAAAATAGAAAAAGAGGTCAATTTCTGAAAACATTGTATAGGTGTAACATAGTTGA
CACTATCAACTACTACAATGGGATCAGGAGACAGAGGGTCAGTGATAACGGAGCTCTCAAGACTGGAACTGGGCGATC
CAGAAGGACAGGCGGTTGATGAAAGAAAATGATGAGTGTGGTTGAGGCTGCGTTTCCCTTGAGGAGGTGCTGGTGCTCA
GGGTAGAGGCATCCATCACGATGTCAGAAATGTGTGTCAGACACGAAGGCAGGAATGAGGCTGTCGAGGAAGCCATGGC
AACATAAACAGAGAAGTGCCACAGCGAGACAACCAGCCAATGAGTTGTGTAGTTTCTTGTAAGTAGCTGAAACAACATA
ACCAAATTGTTCAATTTAGTCTACATATGAAATTTCAATGTCAGAGAAAGGCCTTTAATCTCAAACAGAAATAAACTAA
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GAATCTTACCTGAGAACAAATTTTCATGAGCATCATTTGGAGAGTGGGAGAGCATGAAGACTTTTACCCTTCTCTGCA
GGAGAGAATTGTGCAGGAGGAGGTGGGGAGAATGCCACCAAGAACAGGTTGCCTGGGGCTAGTTTAGCGATGGGGTCCA
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CCAGGTTTTGCTTAAATGCTACTCTTTGGAGAATATCAAGAAATTTAGATGAGTTTACCAAATCCAAGATTGTGTTCCC
TGCTTTTAACTCTTGTTACTGAAATAGCCCCCTGATCCCCAAGAGTAATGCTTGACTGAGGTGTTTGCATGAATTGTTT
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GTAAATTACTTTTTGACTTTGCACAATTATAATTTAATGTCATATATCTGCAAAAGAAAACACTTTTATGTTGCTAATA
TTAATTTCCCTTCATAAATTTGAGACTGTTTTGACTATAGATAAATGTAAAATGTCAATGTGGTGAGAATGACTCAGCT
TCTCAGATATTCATTTATTCTATAAAATATTTTTGAATACCTACTATGTTTCAGGCACCTGATCTAAGTGTTGAATAAGTA
GACAAGGACAGCCTGCCTTCAGGTTTTTTTACATTCAAATAACACAAGATGATGAAGAAATTTTTAAATAATCTGGTTC
CATTTGGCAGATAATATAGATATGCTCAGTTTTTATAAATTTTGATACCTAAAGTATTGTGATAATCCAATCATGACCT
TTAGACATTACACTATGCTTATTGATTGAAGGTTGACATATGTTTAGCATATTCTCTTATAATGTATTTAAGGACTTCA
GTGAGTAGACAAAAGAAGTAAACTTGATCAGACAACATGATTTCTGAGAACAGACTCTTCTTGAGCAAGCATTTCTGGT

[illegible]

Fig. 9.271

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TATGGCCAGTCAGAAAAGAGGATCTTCTTTTACGTCTTCTTTTAAGAAAGAAAAAAATCATTTCAGATGGTCTCCC
CTTGTCCCACCTTCCCTTCAATGTCTTATTGGCCAAAATGGTTTCACATGCCGATGTCTTAACAAATCACTGCCTTGGA
ATCGTACTTTCATGATTAGCTGAGACAATCAGGTTTCCCTCATGGTGGCTGGGGCTGAGGCCACCTCCCTGTAAGTAT
GTGGCTGCAAGGGTAGGTGGATATTTGAATTTCACTGGGGGCCTAGTAGGAAGGAATGGCTCTTGGGTAGGCTACCCAT
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GTTATCTGATCTGAGATATAAAGCTCAAGAGGAGATAGCCAAGGAAAGGGTGGTGGAAAGGAGAGAGAGGGTATCAGGTA
GACACAGTATCATGTGCAAATGACTGGGAATAAGATAAAAATGTGGTAGGTTTGGGGAGATGCAGAAAAATTAGTTGGCC
TAAGGTATAGAGCATAAGTTGGGGAGTCAGTCAGAAATAGGGCTATAGAGGTAACTGTAGATCATAAAAGATCTGATA
AGTCACATAGAAATTTCTACTTTATTCTGCTGCAGTGGGGAGCCAGTGAAATATTTTAAGCAGCAAATGACATAATTAA
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ACATGTGAGAATCAATTACAATATTCCAGGAAAGAGTGGTGAAGGGCTTTAAGATAATGTGAGTGGGGCTGGAGAGAGG
TGAATAGGTTTGATTGATGTTTAGGGGTTTTGACTAAACAGGACTTGGTTAATAGGTAAAGAGAAATAGGGATTAATAAT
CACTCACATTTCTGGCTTGGACAAC TAGGAGGATGGGAATGACATTTACTGAGTTAGGATATCCAAAGAAGAAGCAGAC
TGGGGGCAAGGGGAGTGAATTAATTTAAATTGTAATAATGAAGGGCTTATTCTGTATGAAATTCAAGAGAAGTATTCC
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TCCCAATGTGGCAAGTAGAAATGFTCAATACTTTGTAAGGCCTCAGTTTATAAAGCAGTATCATTGTGCCACATTCT
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ATGCTGCAATGCCTAAAACATAATGTAAATATAGATAGCATTGGATTGACCCATAAGCCATAAAGATTCTAGAATACC
AGTGTTCCTCACTATGGAGTGCCCAATGAGCTTCTTGACAACTCAGGAGAGTAACCTCCATTGTGTGACCTCCTTTC
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AGCACTTGCTCTCCTTTTCATTCTGTGGCTTCTGTGCTGCTAGCAGTGACCAGTGACTCAGGAAGGCTGCATATAGTGAA
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CTAGCTTTCTTTTTATGACTACTTTTCCATTCCACACTTTTCTTAATGTTGCAGGAGATAATTTGAAAGAAAGGAAAAA
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ATTAAATTCTGCTGAGACCCTTAAATATTAGCTGTACAGGGGTTGTGCTTGAAAACTTGATTCCAACCCTAAGCATA
AAAATCTTGTCTAGGCAGGGCAGTAGATTGGAGGATTATCATTACACATTATAGGAGGAAGGCCCTGTTCATGCCAGG
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GGTGCCATGTTGGTGGAACAGGAGCAGAAAGGAAATCAGACTGAGAGAATGAAAACCAATAGAAGCACAATAAACAGA
CTGAACAAAGTACAGATGGAGTCAGTTCAAGTTTAAATATGAATGTCCCTGGGCCCTGTGTTTCATCCCAGTGTTAAGTG
CAGGAAATCATTGGATTTCTCCAAGGATGGGTCTTTAGAGCAACCGACTTTAGAAATTGAATACTTTAAAATTATTTTC
TATAAACAAATTTATCAATAGAAAAAATAGATTACTTAATTGGTTTATCTTAATGAGAAAAACAACAAAAGCTTCAACA
AATAATTGTTGCTGATTGGTTGGTGGGTTCTTGGTGAATAATGAGAAGGAAAGAAAGGAAGAAAAGAAAGAGGAGTA
GAGGAGGAAAGGAGGTGAGAGAGGGAGATTTGAAAGGGAAATAGCGGTGCTCTAAGTTTCACAATTTTAAAAGCCTAGA
GTTATTAACAAATTATGCCCTCAATCAGATTTTATATAACTTTTTTCAATCTTGTCTATATTAATGTGCTGTATTTCAT
GAAATGATTTTGAGATTTTAAAGCAATGATTGACAATATAGTAGTTCAATTTAAAGTTTTTACAAGTTGCTGCTGACAAA
ATATGGGTAATGAATTACATCAATAAGTATAAATATAAGTACACGCTTTGAAGTTAAAACTCAGTAAGTTGTTATGAT
TAAAATTGTTCACTTTATTTTCTCCTGTGTATGGGTTTCTCATAAATGGTAACTTATACCTATGAAAATACAGGGTTC
TATAAAGTCTGAAAACATAGTTATTATATATCTTAACGGACCGAAGATGTCCTTCTGATATTATGTATAGCCTGTGTT
TCCAGACTAAGTGCCAGGCACTATTCTAAGTTAGGCACGATTCTAAGTGTTTTACTGTTCAATTTAGGCTGTCTGGCTAG
TGTTTTTCTCACCTTATCAGCATCCCATTTTTATTAAAGTGAATTTGCCATAACACACCTGCACACGCACATGCATGCGC
GTGCACACACACACACTTTTTGCATTTGGAAGCCTGGCTATTATGGAACCTTAGAACATGAGAGCTCAGGGTCAACCAC
CAAGACTCAATAAGGTATGGCTGGGGATGTGACCAAGACTATCCTAGCATCTATGGCTGTGGACTGGGTGTCTTCTCCA
CTGACCCTTGAAGGTAGTTTGTCTTCACTGCTTTAGAAATGGTGCAGAGACTTAATCTTATTTCACTTTTTCGGTGTGTTG
TCTTGGTTCTGTAGAACTGGCTTGAGGCTTAGAAGATGTCCTCCACCTGCATTTGGAAAAATACATTTCTAACACTT
CTCCTCATATGGAAATTTTTAAGTCATTGAAAACTCATACTGCAGCATTTGTAGAAACAATTTCAGAACAGAGTACCT

Fig. 9.272

GCAGTGCACAGGCGGGCGGGAGATTCTCCAGGGATCCTCCCTCTTATCTGCCTCCTGCATCTATCACCTGCATTATTGG
GAACTTGGTTTGAATTGTTATATCCTGGAGAAGACAAATTTATCTAGTAGGTAAAAGGCAGGGGCCAAAAAGAAGTA
GACGTATGGAGACAGCTGGTTCGAGCAGGGGAGGAACCATGTGAGCAATGGGAGAGCAGATTTTACAGAGTTCAGATG
GAATCTGCTGAGGGATTTTCTTTCCAGAGTCATGAAGCCACTTAGCGTGGTCACAGATTCTCTTTATATGTTCTCTA
CTTGCCCACTATTGTTGACACAAGTGCAATAGGGTTTATTGATGACAGCACAGACGACTCTGTTCTGCAAGGAGGTAAT
CTAGAGCTAGTCTGTTGTCCATGACTGCATTAGCTACTGAATTCAGTGAAGCCTTGAGTTTAGAGATTCTGTTCCAGT
TCTCTGACCTAGGTCTCCAATTTGTGTAGAGAGGTTTGGAGGGTGACTTTATGATAACTGAATCCTCTGTATGGGGCA
GCTAATCCTATGGCAGCCCCCTAGCCCTACCATTATAAGGCCAAGAGCTCTCCTTTTTTAAGAATTTGTAATGTTATAGA
TGGTTACACCTGTTCTTGTGGCCCCAGGAGTCTAGGGCACATCCCCTGTATGTTGGGTGTTATCTATGCAGATAAAA
AGGTTTACAAGTGAGGGGGTTGAATAAGTTCTTTTAGAACTTCCCTCACAGGGCAGTTTATTTTGTGTTAGGTCCATATA
AAAATACATAGCCTGGAGGGGTGTAGGCCCACTCGGGAATGTTGGTTTTGAACCAAGTGTAAGCATAAGCAATCCCACC
ATGAATAAGATCCAGAGTACTGGGGCTACCTTATGCTGAGCTAGCATTAGTTTGAAAGAAAAGAAGGCATTACTTATAA
TTTCATTGAAAGAGATATTATAGATCTTCCAATGACTCACAGGAATACTCAGGTGCTGGTCCCTATTGTTGATTGTTTCAG
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GAAAACAGTAAAAGGGCCTTTCCATGAGGGGATTAGTTGAGATTTTGTAGATCCATCACTTTAGGTTTTAATAAGGACC
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AATTCTCTGTCTTGGATTTGGATACTTTGTACCTTACATTTTGTCTAGGAAGKTTAAGGTTTGTACTGCATATTGTTGTG
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TAGATTTCTTTGCTGGGTCTTGTCCAAATAAGTGTGGGGCATCTCTAAAACCCTGAGGTAGGACTGCTGGCTTTCTCTGG
GAGAACTGGGATATTAGGGGGAAATATCAGTCCAGATGTTGGGTAAATATTAAGTGGATACCCATTTTGGAAAGTATA
TTCTACTCAAAAGCAGTGTGAGGCATTTAGACATTGCCAGGGACTAGTGGGAGAATAGTAATTAGTCCCATAAGCAAT
ATAAATAAAGGGGATGTGAATCTTTAGAGGAAGGGGTGCTACTTGCTCCTGTTACCCAACAGAATTTGGGGGTAAGTT
GCCTAGAGAAAAAGGTTAGCACAAAGTAGGCAGTTCTTGTATTTAAAAGGATATTTATAGCACTACCTGTCAATTTCCAG
AGTTTCCCTTGGCTCTGTTCTTTAATGATGATGTCTGATTTGGAAGCTGGCCAGAGTGGAGGGGCCCTTCAGCTCAAG
GCCATTATTGGATGGGGGTCAACTGGGGGACCTTTTGGATCCCAGGGCAGTCCCTCTTCCAGTGGGACTTATCTCCAG
CTTATCACAAAGAGGGCAGGCTGTGTGGGGCTTCTCTCCATTTGGCCCATTTGGAAAATTCCTTCTCCAGTAGCCTGGT
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GACTGAGGAGGCTAATTTGAGGATTTGTGGCATAGGGGCACTGGGTCTAAGGCTGACTTTTGTAAATTTTCTTAATGT
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TTCCAAATTAATCTAAATGTAAGAGCCAATTTGGGGGAAGCCCTCTATGAGCCTTTTGGGGTCTATCTGAGAATTTT
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CTACAAGCTTGGAGAGGAGAACTAAGGAGCCCGGTGTAAAGTGGGGCCAATTCCAAAGGTCTGGGAGTGGGATAAATAG
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AAGAAATGTTGGAGAGGAATCAGCAGAGGAGTCTCCAGGGCCGATGCAGGGCCCTGGAGGAGATGGCTAAGCATAGCTA
GGATGTGAGGTTTGCCTTACAAGCTTTACGAGGATTAGGGTCTTCCCTTTAGGCCATAAAGGTTTGTAATAGGGGACC
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AGACTTCTCTGCAATGAAAAATTAGTTTTTTTCAATTTTAGGGTTTGGTGGTTGGATTATCGCAGATTTTAAGGATGCA
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GAGTCCCTTCTTCTTCATCTTTTTTCTTTTTTCTGTTTTTTTTTTTTTTTTTTTTTTTTTTTGGAGACAGAGTCTCTCTCT
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CAGCCTTCCGAGGAGCTGGGACTACAGATGCATACCACCAAGCCCAGCTAATTTTTTGTATTTTGTAGTAGAGACGGGGTT
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GAAAGCATTCTTCTGATCCCTTATACCTTGGATCAGAGTAGTGAGTAGGGCATCCCCCATTCATCTCTGGAGTTCTG
GAATAAACCAGTGATTACCAGGTACCACTAACCTGGTCCCACATTTCCCTCCAGGACCAGCCTTCATCTCTCTGCTAA
TGGTAATCTGTCTGCACTGTGGCCTCTGGCTGACCTGTACCTTTGACACTGTGATCTTACAGTGACCTTTGTTTGG
GCATTTCCAGCAATGAAATGGTTTTCTTTCTCTCAAATTTCCATTCTCCATATCCCTTTAGGTAGGTGAGGATCTTATC
CCTAGCCAGCAGCTGTAAGGGAACCTGGGCCTTCTTTCTCTGGACATAGCACTGTAGGTCCAGTATATGTTAAGAATG
TAGATGGTCAGAGGAACAGAGGAAAACCTGCATCTGAGTCCCCTTGTCTATCCCTCCTGTGGATTCTTGGGGGAAGCATGG
AAAACAAGTATTTAAAATGACAGGACAATCCATCTGCCACCCGTGGGAATGGTAGAAAATAAGGGATATTTCATGGAAGG
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GAACTGTGAGGGCCAGGGAGTTTGGGATGAGAGACCACAAAAGGCAGAGAAAGAATTGTCCTCTCCCCAAAGTGCAGAT
AGCTCTAGAAAGGAAGTAGGAGTTGCTCTTAATGGACCACATACAGATGCCCTATGGAGACAAACAAATGACCTCAAGA
GCCACTGGAAAACCTGGTCTCAATGTGTAGCAGGATTTAAAAATTCATGATAGGAAATAAAGAATTGGTGGAGACAGAG
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CAGACAGTGTGGCCAGCATATAAGGCCATCTCAAAGTCCACAGAAAAAAGGAAGCATAATAGGGTGTAGACTTATTGG
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GCCACCTGTCCAAAAGGGCCACAGTGACTTAGGTCTACTCAGTACAGACTCTGAAGTCCTCCACCTCTGCTGTCGCCCCA
CCAGGATGAACTGAGGAATCTGCTAGAGGGAAGAGTGACCAAAGAGAATTTTTCTGGAGATGGGCTGGTGAGTGAGACA
GCAAGAGAAAGAAGACCTCACGTAAGCAGAGTTGGGTGCCTCCAGCCAAAGATGGCAAGGCACAGAGGGTCTTACCGAG
AAGCTGCAATCTGATTCATGACACCAAATGTTACTGGCAGCGGGTTTGGGCAGGATACACAGTCTTTGGTTCTTATTG
TCTGAGAAGAAAAAATACATCCAAGAGACAGAAGTAGATTTAAGATGGCAGACAGGAGGCAGGACTAGATTGCAGCTCT
GGACAGAGCAGCATGTGGAGGCTCGCATTGTGAATTATAGCTCCAGATTGACTGCAAGAACAACCAGCAACCTTGAGA
GGACCCACACACCCTCTGAAGGAAGCAGACTGCTCTTGACAGGACCTGGGAAACACCCCCAAATACTGTGAGTACCCCAAC
TGTGGAAGTGGGAAAGGGAGACCCTCCTCTCCTGAACACACACCCCCACTGGAGAAGCTGAAGGTCTGTTTGCAAGAGA
AGTTTCTGACTTTACCTGGAGCTGAGTCAATGTGGAGAGCTGAGTGAAATACAGAGGCAAAGAAAGTAGCAGAAAGGCC
CTGGAAGCTCTCTGAGTCCCCTAGCAGGCCATCCATGCCTGGCATCACAGGGATCCATCCAGAGGGCAGCCAGAGCCAG
AGGTGCAAGGGGTAAACTCTACAGGGAGAAGAAAATCTCTAGCTGAAGTTTGTAAACAATTTGAATGGGGTGAGAAGCC
TCCTGGCCAGAACTCAGGGGAGGGCACACATCTGGCGTGCAGACTCCACAGGTGGAGTAGAACCAATCCCTTTCATTCA
CAGCTGGGAGGTGGGTAGCCTGGGGCAGATTTTCAAGCTCATCTTGCCCTCCAAGTGAAGTGGACTCAGGCTGTTAGA
GGGTGGGAGACACAGTGAGAGTGAGACTAGCCATTTGGTTTGGGTTTGGCTGGAAGCAGAGTGAGGCCTGTGACTGCTG
GCTTTCCCCCACTTCCCTGACAACCTGCATGACTCAGCAGAGGAAGCCATAATCCTCCTAGGTGCACAACTCCAGTGAC
CTGGGAATCTCACCCCCATCCCCCATAGCAGCCCTAGCAAGACTCACCCAAGGAGAGTCTGAGCTCAGACACACCTAGC
CTTGCCCCCACCTGATGGTCTTCCCTATCTACCCTGGGAGTGGAAGACAAAGGGCATATAATCTTGGGAGTTCTAGGG
TTCCTCCCCATACTACCACAGCTGATGCTCTCTGGAAGGCAACACCTCCTGGCAGGAGACCAGCCAGCACAAAAATAGA
GCATTAAACCACGAAAGCTAAGAACCCCCACAGAGCCCATTGCGCCCCCAACCCCTACCCCTGCACCAGAACAGGCA
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GCCCTATCTTCAGCCTCCTCAAACAAAACAATTATCAGCCAAGAATTTTGTACCCAGTGAAATTAAGCATCATATATGA
AGGAAAGATACAGTCTTTTTTTCAGAAAAACAATGCTGAGAAAATTTGCCATTACCAAGCCACCCTACAAGAACTGCTA
GAAGGAGCTCTAAATCTTGAACAAATCCTGGAAACACAACAAAACAGAACCTCTTTTTTTTTTAAATTTATTATTATTA
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GATTTCCAATTTTCATCCATGTCCCTACAAAGGACATGAACTCATTTTTTTATGGCTGCATAGCTTTCATGGTGTAT
ATGTGCCACATTTTCTTAATCCAGTCTATCATTGTTGGACATTTGGATTGGTTCCAAGTCTTGTGCTATTGTGAATAGTG
CCGCAATAAACATACGTGTGCATGTGTCTTTATAGCAGCATGATTTATAGTCCTTTGGGTATATACCCGGTAATGGGAT
GGCTGGGTCAAATGGTATTTCTAGTTCTAGATCCCTGAGGAATCGCCACACTGACTTCCACAATGGTTGAAGTGAAGTGA
CAGTCCCACCAACAGTGTAAGTATTCTTATTTCTCCACATCCTCTCCAGCACCTGTTGTTTCTGACTTTTGAATGA
TTGCCATTCTAACTGATGTGAGATGGTATCTCATTGTGGTTTTGATTTGCATTTCTCTGATGGCCAGTGATGGTGAGCA
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GAACCGTTTATTAATAGGGAATCCTTTCCCCATTGCTTGTTTTTCTCAGGTTTGTCAAAGATCAGATAGTTGTAGATA
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CAAAACAAACTTTAAAGCAATAGCAGTTAAAGAGAGACAAAGAGGGATATTATATAATGGTAAAAGGCCTTCTCCAACAG
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CTTCAAAACCATGCAAATACATGGAAATTAATAACCTGCTCCTGAATGAGCATTTGTGTCAAAAATGAAATCAAGATGG
AAATTATACAATTATTTGAACTGAACAACAATAATGACACAACCTTATCAAAACCTCTGGGATACAGCAAAGGTGGTGCT
AAGAGGAAAGTTTCATAGCCCTAAATGCCTACATCAAAAAGACTGAAAGAGCAAAAAGACAATCTACAGTCACACCTCA
GGGATCTAGAAACAAGAACAACCAACCCAAACCCAGCAGAAAGAAAGGAAATAATCAAGATCAGAGCAGAACTAAATG
AAATTGAAACAAAAAAAACCATACAAAAAATAAATAAATAAATGAAACAAAACCTGGTTCTTTGAGAAAATAAATAA

Fig. 9.275

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AATTGATAGACCATTAGCAAGATTAACCAAGAAAAGAAGAGAGAAAATCCAAATAACTTCACTAAGAAATGAAACAGGA
GATATTACAACCTGACACCACTGAAATACAAAAGATATTCAAGGCTACTATGAACACCTTTATGCACATAAACTAGAAAA
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CAGCATTCTACCAGACATTCAAAGAATTGGTACCAATCCTTTTGACACTATTCCACAAGATAGAGAAAGAAGGAACCTT
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CCACATAAACAGAATTAAAAACAAAATTCATGATCATCTCAATAGATGCAGAAAAAGCATTCACAAAATCCAGCAT
CCCTTTATGATTAAAGCTCTCAGCAAAATCAGCATAAAGGGACATACATTAATGTAATAAAAACTATCTATGACAAAC
CCACAGCCAACGTAATACTGAATGGGGAAAAGTTGAAAGAATTCCTCTGAGAACTGGAACAAGACAATGATGCCCACT
CTCACCCTCTTCTTCAACATAGTAATGGAAGTCCTAGCAAGAGCAATCAGACAAGAGGGAGAAATAAAGGGCATCCAA
ATCGGTAAAGAGGAAGTCAAACCTGTCCTGTTGCTGATGATATGATTATTTACCTTGAAAACCTCTAAGAACTCCTCCA
GCAAGCTCCTAGAACTGATAAATGAATTCAAGAAAGTTTCTGGATACAAGATTAATGTACACAATCAGTAGCTCTTCT
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CCCCCAAAGGAAGGTCACATATCAGGCAAACCTCTGACGTTTTTGCCCCCTTCTTGAGCATGCCTGGACATATCCCCGAAGG
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CGTTGGAACCTGAATTACAGGACACCCAGCTAGTGTGGCTGCTTGGTGTGGGGGGGAAACCTCACATGTTTGTTCGTAG
AAGGCTTCATCTGTGTTGATGATTTTTGTGGTGTGAGAGTAGAGGAAAAATGCCATCAGGGAGAGTTTTCTCTACACCC
TATAGCTCCTAGATGCTTTTTTATTAAAGACATTCAAACCTTGTGAAAAACAAACAAACAAAAATCTTTTTATTAAAG

Fig. 9.276

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ACGGTCAAACCTTGTTCAAAACAGCACAGACATTAACCTCGGAATTTAGGATTTATTATTGATTAATTGACATTATCTTC
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GATTCCTCCTGCCTCAGCCTCCCAAGTAGCCGGGACTACAGGCACGGGACACCACGCCTGGCTAATTTTTTTTGTATTTTT
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CCCCATGGCGTTTTATTTCAGGTGTATGTATGCCAGAGATGAAAAAGGCAAATGACAGAAGAGCAAGTATATGGAGA
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AACAAACCTAATCATTCCTCTCATTTTTGATCAGCTTTCATGTATATTTTTCTAGGCCTAATCAAATCTTCTTTGGTGTG
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CAAGGACCCATAAGTATTTATTGATATATCTTCTTAAGTATGTGATAATATTCTAAATAGATAGTAGATTTAATCATTG
GGATTTATGTGAAGTTGGATTAAATAGATAACATGATAATTACATTTAGTATTGCATGGAAGGACACATTTTCTCACTT
ATAGAAGCCTAAACAAATGTCACAATGTTGATAGATTCTTTTATAGGATGTTAACTCTGAAGCTGTCGTGAAAATGTGT
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TGATCCCTCCTATTTCAATCTCATAAGACACTTATCTCTCCTCCAGAAAGATCTGAAGGGTTAAGTCACCTATTTATAC
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CAGTGCCTAACAATGACACAGGATGACTCAGTTAAGGAGAAAAATACTCCATTTATCAGCACAGAATTCTTCCCTTTAA
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TTATTATCCTGACACTGAGGACAAATATGAGTTGCTGTCCATCAGCATTCCAAAGCTGTTTTTTTTTTTTTTTGTTTTTTTTT
TCTTACACTTAGCCTACTTCACTCCCTAACACCACAGCCTGATTTCTGAGGGGGCCTGAGGTAGGAGGAGGAGAATGG
AGAGCACTTTCTCTGAGAGCCGCCATTTTAAACAGATCATTAAAGACACGATATTCACATGACGGTTGCTTACTCTCTGA

Fig. 9.277

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TGAAAAC TACAAAA CAGAATA CACAGGGA AGGTAAT CTGAAGGT GATACCT TTTTCCT ATGATCCT TGGCCTT ATAAAC
CACTAAT CAAGCCT GAGGGCC GAAGTTC CTGCCTC ATCTCTG CCTGATG TTACGAAC TAGGCAGG CTAGCAG AGGCAAC
AGGCAAC CTGGGCAC CTGAAGAG CTACCTG GATAAAC CTAGAAGA CAGTAGG AGGTTAA AGATGAGG ACAAGTCT ATCA
AAACAAA AGCCTGT CAAGACC AGAAAG AGAAAGT CACACTT TTTGACT TTTACAG TTTGTG CTGGGCT GACACAA AGGCCT
CAGTTAA CACCAA ATACAAC TTTCCAT AGACTTT CAAGTTT CCCTCAT ATTTTTC CTGGGTC ACTATTCC AGAGTTG AGAA
ACTGAA CTACATCT AATATTT TATTAAA ATAAAT TGAAGAT TATTAGA AAAAGT TATTAG TCAATAG TAAAAC GTTAT
TTGGT TAGCTTT CTAAACA ATGGCAT TTATAGA TATCTG ATTATCC CAACAAA ATCTTCA AATGGT TCATGG ACGCT
TTGTCA AGCTTT TGTGCC ACCTG AGAGAAA AAAAGATA AATGGGG AGGTATA ATGTTA ATTTTGT AGCCTTT GCTTAAT
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ATTTGA AAGCAT CCAGA AGAGACA ATCCTT ACCACAG TCTCTTT TAAACT CTTAAT GGCTGT CACAAA ATTCTAT TTTT
CATTTT CTCTA AGAGCAT TCTAT TAAACT GTTCCTA GTTATT AGCTTT CATAA AGGCAC ACAGAAA ATGTTT CCTCTAC
AGCCACA AATATT CCTAA AGCAGA ATCATAG TAAAAC AGCTGT GATAAT GTTTTAT TCAATTC CAGAAT CCTAAT GATTG
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GCACTAT TTAATG TAATTCT CCCAGC CACTCTG ATTTCA CTCAAT TTTACA ATTGAGG AGACTG AGGCAT AGAAACT AA
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GACACT GCTTTT CTGTGT AGCCC AGAGTGT CAGCCTC AGTGCTT CAACTT GAGCTTT CAGGAT CTATTTA AGATGGA
AAATATA GTTACAT TATGTC ACCATTT GAGATG CAGAAA AGACAGG GCCACCAT TTTTGC ACAATTC TGAAAG CACAATTC
ATGCTGT GGT TTTGTAAA ATGGTACT CCCTAG AGTTGGG CAATGG ACAGCTC ACACAG AGATGC AGTGGC CCTGCTAAA
CTAATGC CTTACATA AAAAGAG TTTTACT CTTCAAT TCCCTT AAATGATT CTTAGT AACACT GGGCTG ACACAC ACACAAA
CAGTTTCT AAGTAG GTCTGT CCGCATA CTATGAT CACTGGT CAACATTA AGTTTCC ACAATTA ATTTTAC ATAATCCA
AGACTGCC AAGCAG TTTATCT GGGTAA ACTAAAC ATTCTGT AGTCAT TTTATCT GCTTCC AGTCAT GCCCAG CAAACAGG
TTGAGA AGACAA ATGTTCT CAGAAAT GATCTCC AAGGAG TTGGG AGCAGG CTGCTTAT ACGTCTA ATTCACC AGAATAG
GTGAGCG TGGTCT GTGGT GACCTTTT CATACT GTTGC AGAGCTG AGTATGA AGAGAT GACTCAC AGTCTT CCAATG CAAC
ACAGGTG ACCCTG CAGCCAG ATTTTGC CTTCA GTGGTAT GTGACT CCCATG GGGGT CAGGAG AGTATCTC AGACATTGAA
ATTTGAC CTAATGG CCTACCT ACCTGC ACACGC ACCCTG CCCACTT ACAGA AGGGCA AGAACTCT GCATTACT GAGGCC
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CAGACTCT ATCTTAG TTCCTA AGCCACT CACCAG GTATTT AAAAGA ATGATTTA ACACA ACTAGA ATCATTTA AATAAC
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CTCATGT GTTCAG TTATTCC AGAGAA AACATTA AGGAGA ATTGTATT CTCTTCC CAGCTAA ATTTT TAGGTCTT CAAAG
CTGGCA ACCACA ATTTAT GCTTTT CTA AAAATCAT CTATG ATACTA AGTATG AGCTGG ATCATGA ATAAAC CCTTGTAA
CAATAA ATGCTT GATGACT TACTTC ATCTC ATTAG CAAGGGA AGGTAA CTATAA TTATCA AGGTACT ACAGGAA ATAG
GGCACTAT CTGGAA ATGCTAA ATACAT CTCTTGC ATGAAT CACTTCTT ACCAGA ATTCCCTT TCTTTCC AGATCCCAC
CTCCAGG GTTCTGT GGTCCATA CCAGGC ACTACAT TCCCTC AGGGGCTAC AGCCTATG AGCGTCTCAT GGGGCTATGAAA
CCATTTG AGACCTG GAAATGA AGTAATTG AATACAA ACAGAAA ACTGCAA AATGAGG ACTAAC ATTTA ATTAATGTT
CAAAGC ATAAGATT ATGTCG ACTTCA ATAATTGT CAAATG AGTATTCTT AACATTTT TACTAA ATTAAAAA ACTTATGT
GCTGAG TTTTTT ATTTTACA AGTATCTCCA AGTATGCTGG ATGATTG CAAAGAAA ATCAAGGCC AGTCATTGG TTAAAT
GAGTTTA ATAGTAG CCACATA ATTTCAA AAGCAAA ATTATAA AGACCTTCCC CAGACTGT TGATAG CAAAAA ATAATCTA
CGTTGTG GAAAGTGG GTCCAT GTTAATAT GTTAGA TATAAG TAGTGAG GCCTAAA AAGGTATT AAAACATCTTTGCTTA
AGGTACT ACCTATTT GCAAGATT GTTATTTT AAAAATAG CTTATG TTTTAA ATTGTTATT GCTTTT TATCACTCTAATA
AGAATTT ATAGTTG CTGTA AGATAACA AGAAAA AGGTAA CTATCTG CAGAGATGC CTGAGAGT CAGCCAGG GAGTAAC
TTAAATC CTGCAG GAATCTG ACTCACT GAAGACTGT CAACTGA ATGAAGG CTTAA ATTTT CATGGTCTTGGGTGGGAGAA
CTTTTTC ATATTCTT TCCCCA TATGGA ATAACA ATCTGCC CTGAAA ACAGGG AGTATTTTGG CATGATCTCTTTTGTCT
TATTTGC CTTCCAT TTTCCATA AAGCAACT TTTTGCCA AGCACCATA CTTAAG ACTCAACT TTTTGTG CAAAAA ATATCAGA
CAAAGC ACTGTCTTTA AGAACAC AGAGAAC ACACACTAG ATCCCTTCTTCTG AAAATCA CTGTTCTAT GTTGTGTTGTGGAT
ATTTT TTTTAGC ATTCACT GCATGCC TGGAA TGAATAGG CTGTGTTTCT CCCCCA AAAGAG CACAAATTA ATATACA AGGT
CAGGT TAAATAGTTA AGTCTGCTTTCCT ATCCCTTATACAGA AGCTATCC ATTTCA AAAAAAAAAAAAAAAAAA ACTTCTTT
TTACTTT TTTTAA TTTTAA TTTTGGTGC ATACATA GTAGGTGT ATATAT TTTGTGGA ATACATA CAAACATA CAATGCA
TAATAAT CACATCAT GGTAAATGGGTATGC ATCACCTCA AGCATTTAT CCTTTT TATGTTAC AAACAATCCA ATTTT
ACCCTATA ATTGCTTAAAAA ACATACAATAA ATTTGTTAA CTATAGTCA CCCTGTTGTACTATCAA ATATTAGCTCTTAT
TGATTCT ATCTAATTAC ATTTT TTTTATA CCGCTA AGTATCCC ACCCTTCCCTCCACGCTACCCTACCTAACCTCCGGA
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TCTTTCTTTTGGGTATGTA CTTAGCAGTGGGATTGCTAGATCATATGGTAGCTCTATTTT TAGTCTTTT GAGGAGTCTT
CAAAC TGTCTCCATAGTGGTTGTACTAATTTACAT TCCCATCAA AAGTGTACCAGG GTTCCCTTTTCTTTACATCCTC
ACCAGCATTTCTTATTATTGTCTTTTGCATAAAAAGCCATTTTAGCTGGAGTGAGATGACATCTCACAATAGTTTGTAT

Fig. 9.278

Fig. 9.279

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ATGCATCTTTACCAAAAAGAGAGGTAAGAAAGGAAGGATAATCCTGCCAGAGATACATATGCAAAGATCTGAAAGTA
ATCGAGAGTGAGGGGTTGCAGATTGGACAACACTTAGTAGTGGCTCCAGATAAGGTTCTTAAAGACTGGCCAGGATGCA
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CATTTTTTTCTTTTAAAGTCACTCAAACCAATAATTAGTACAAAAACAGAAAGGTACATTTTCATCTTTGAGGAACT
GGGAGATGTCTGTGACTCTAAGCCACAATACATAAAGATAGAAGTAGGAAGAGAAAAGGTCAATGACTTAACCAATCA

Fig. 9.280

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GAGAACAATACTAAAGTGAAGCAAACCTGAGCTGCCCCAGTCGCCCAAACATACTTTGGAAGATGGAAGAGTTCTAAAT
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TGCATTTTTTTTTTAAAAAAAATAAGGTTTAGAAAACATTTGTGCAAGGGAACAGCCTTAATACGTGCAGCCACTTGTCCA
GGAATAATACATTCAATTTTTTCAAGTTTAAATTTCCAGTATGTTCTGATCCAAGGGTGCCTGTTACACTCTGCTGAATTT
TTAAGAGGTAATTTACATCTCTACAACCAACTCCAAGCATGACATTTTATTACATCCGCTCAAATGAACAGCTGCTA
AGTCATCAAGTTCTCTCAACTTTGCTTCTAAAGAGAAAAGTTTAGTTTTAGTGCTTCAGTGAAATACTTTTTTGAAGA

Fig. 9.281

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ATGACTTTTATAAAAATTCATTTGTTTATTATGGTAAATAAACAACTTAATGGCAAGGGGTGTGTTCCCTTGTAAGGTGA
TCACGGAATGTTACCCAGAATAACACAGACATCTCATTTCCAGAGAAGGAAATGGTTTTATAAGTTTGTGAGGTCTC
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CTGGATTCTAAGCAGGGACACACAGAAATTACTTTCGCAGGTAAATCAGCCCACCCAGCCAAAGTGTGGAGAGATTTGT
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GAAAGTTTAAATATTTAAATATATACCTACTATATGTATTAGCAAGCTCATCTTTATATATGCCAATATCTACTTAGAC
ATTAAACCACAATTTGCTCAGGTTATATAATATGCAAATGAGTATTTTCACTGTTGGTATGATAAATTTCTGAATTTT
CTTGCAACTCTGACCAATGGAGGATTATAATTTTTTTTATTAGCTCTTTTCTCTTAAGGTGTAAAAATCTAGACTGGCT

Fig. 9.282

TAGCATTCTTATTTTGGCCTCATGATCACAGTGGAAAGCATTTCATATAAGCTAAAATGTCCAATTTTATAATTGAGAAT
CAAAAAGATGGAATAACATTAATATGTTGGAAGGAAAATAACCCATTAACCCCATTTATTCATCCTCAGTTATTCTCAT
TTTCAACTTACCTTACATATCTTAAATAGCTGTAAATACTATTTCTGTACTCTGCTTCTTCATTTAACATATCAATATT
TTTCTATTTTGGCCAGGCGGCGTAGCTTGCTCCTGTAATACCTGCAATTTAGGAGGCTGAGGCAGGCAGATCGCCTGA
GGCCAGGAGTTCAAGACCAGCCTGGCCAAAATGGCGCAACCCTGTCTCTACTAAAAATCCAAAATTTAGCCAGAGTGGT
GGAACATGCCTGTAATCTCAGCTTCTTGGGAGGCTGAGGCATGAGAATTGCTTCAACCCAGGAGGTGGAGGTTACAGTG
AGCCAAGATCGTGCCACTGCACTCCAGCCTGGGTGACAGAGCAAGACTCTGTCTCAATTTAAAAAATAATCCAT
TTTTCTAATACTTATAGCTTTCATTTTATTGAGTGTCTTATGAAAGAGCTCTATCAAGAATTTTACATATATATTATT
AGTTCTTATTACAATATTGAAAAGAAGGAGTTGTTAATCCTAATTTTACAGGTGAGAAAATGATTCCGTGAGATGAAAT
AACTTGAGCAAGGTCATAGGCTGGGGTAACCCAAGTCTGTCTACCTCAAAGAACGTGCTCTGGTTCTTCAGCTTGTTGC
CTTCTCTACTGTGGATATAATTATCTTTTTAATGGCCATAAAAGTGTTTCATGTTGTGAATATTTCTAGATGCATTACA
GTATTTGGCAGTTTCTTTTTCGGAAGTATGGTAGCCTATGAAATTCAGGGTTGACGTTGTTGGTTGTTGAAAAATAGCT
TTGTGAAAGACTGTTGGTTGTTGAAAAATAGCTTTGTGAAAGGCTCTTGACGCTTGAGACCATTATTTCTCTACTGG
TAGCAAATAAGGTCTCTGGTCTTTATAACAGACAATAAAGAACAAGGCCAATGCCTCACAATTTTGAATAGATTAT
CTTTCCAATTTCTATTACCAAGTTTTTCTGTTCTCTGCAAGCTCTGTAAGGCTTCAGGTTGGAACAAGTAGTACATATC
AGATTTTGTATGACATCTAAGATCAGCTTTGAAAAAATATGCAATCTTCCTTCTATTTCTTCTATCTTAGAACTTG
TAAATACGCAAACAATGGAATTTCTCATATTTTCTGCCTGTGCAATATATATTATGATTTGGTTTAGGTATGTGACTAC
TGTCAAAGTTGGTTCAGGGTTTTGAGCATGAGGTAACTTTTTTTGGTCTTATTACCCAAGTTAGTGATGTATCTGTTA
TGAAATATACTTAAAGATATACAGCTTTTAGAAATGATTAAGATACCTTGAAATATTTGATGTATTTAAACATAACTA
GAAAGACACAGAGTGGATCACAGATAGGTTTACTCTTTAAAGTCATGAAGTATGAGATTCCTTGGAAGGCCATTTGT
CCACTGTACTTCTAACATGCTCTGTGTCCCTGGAGGAGTCACCTAGTCCCTCTGAAGCCAAGTCTACTGTCCTGTAAAA
TAAAAGGGGCAAGGCAGGTCAATGGATTTTAAACACGTTTATTTCAATGGGACATTTTTTTTATTAATGAAATGTGAGTTG
AAGTTAATATAAATGAATATTTATACTGTAATTTTGTAGATATAAAATCATATATAAATTCATATATAGATATAATTTT
TAAAAACATAGATATGTGTGAAATATATAAGTAAAAAGTATAAGCACACATAAAACAATAAAACAGCCACTGAGTTTGT
ACATTTACCCCCCACCACAAATTCAAACCCTCTCCTTAAATACTCTAAAGACTTTAGGATTCGAAGGAATATTGTTT
GAAAACCACTAAGCTAATGACTGTTAAGATCACTTACAGTTTTACTCTCTGCAGTGCTGCCTTACCGCGGGAGAGTCAT
TTGTGTATTCTGTATCTCTATATGGATTGGGATAAACTCGTAGGTAATGTAGCAGTTCAAATACCTCTGCCTTATAAGA
CCTTGAAAGTTATGTAACTTTTCCCACTTCTGAGATATCACAGAAGAGTTATCACTATAATAGTTAAGATCTATTCATTT
TATTTAGCATCTTATTTTTTACAAGTTTATTTTAGGCTCATCTGGAATGAGAAAATCATGCCTAAACACTGTATGATG
CATAGCTGATAATTCAGGAAGTGAGTATGGTTGTTGGTAAAGAAATGCTGAATTTGGTATGAAATTTCTTCAGAAGCAAGA
CGCTTCTTTTGGTATTTCAAAATGGGGTCTCCACCCTCTTTTTCTATCATGTCAACTGGCTGTGGTTATGTGTGCAAT
AATACTTAGTCCTGACTGCAACTCTTTTTTTTTTCTTCCCCACGGATCTGAGGCACACCATGGTTTTTGGCCTTATCATT
AAAACCTTGCTTTTAGGAGAGGTAAAGGAATAGCTCACTTAGTAGTTTCTCTGTCAATCATTCCAATCATGTAGTGAAT
TCTATTGAATCAGAAGGGATGATGACATGAAGGTAACCATGAGTATACGAATGGTTAGTCATGGATCCACCCAACCAT
ACTAGCAAGATTGAGAAAAGTAGATTCTAAGATGAACTGAAGGGGAAGAATATTAATGAGGCCTTTAAGGAGTTTTTA
AGCTTTTTTAGGGAAAGATGCTGAGCGATATGTATAAATATTCCCAATAATACTAATACTAAACATATAGAATTGTCAAA
GTTTAGATAAAGGTATAGAGAATAGGGCCTGAGGGATATTAGCTAGGTTGGTAAAGGAAGATAGTCACAAAGAAAGCACA
GTCTTAATAAAAAATCAATATAGATTTTCTTGGAAGCCTTTCCAGAGGTGATCCAATATCTCAAAATATAATACAGGGG
TTGAAGGTATATTAGAAAATGGACAGAAGTAAGGTATACTTCAGCCACAAGCACCAATCTCTCCAGGGATTGGTGCCT
ATGGGAAGGGCTGGGGTACCACATGGGGCACAATGACAGCAAGCCTGGGTGAGACACTGTTAAATATAACTGTGTACAG
GTAACATACTAATGAGAGCCCGTGCCAGTGATATGTCTCAGCTCTCAAATTGCTTAGTTCTTTGTCAAGTTTGACCTAA
ATTAGAAAGCATTTCAAAGATATGGAAAGTAGGAAGAATGCCAGAACCCTATGGATGCAGTTTTTTTAGCATGACAGGGGC
AATATTGTAAGTATTTATTATCTGTCTAAGCCTTCTAAGGGGATTGGCTCCCTAATGAGCATCCTGGGTGATGGCTGA
GGATGCGAATTAGAGAAGCTAACAGAAGGGTGAAGTGTCTTTCAGAATAAGAGTTAGAAAAACACAAGGGAATATTTCA
AGAGAAGAGCTGGGGAAGTGTGGCAAGAAAAAGAGATTATACAGAGACCAATATGGAAGAATTCAGCCCAAGTTTGAA
CTAAGCATCCACCATCTAGCACCTTACCCTGTTTAATAACACTGGCAACCTCTGGGCAAAAGCTGCCATTAATGAACT
CAGTGATAGAGGACATTTGCCCCCATCTCAGCCCTAAGAATTTCTGCCATAGGGTAAAGGGGGTACCTGCAAGCCCTT
CCATTTCCCTACTACACATGAGGAAGAATAAGAAATTAGAAAACCTAGAACTAGGTGTAGAATGTATGCACACCAAGGA
TTGCATCTTCCAAAGTACCCTTATAGGAAGCAGTCTGTGTTTAAACCCACAGAGTTAGGAAGTTTTCTTTATATCT
AATACAAATCTTCTTTGCTTAAATTCTAACCCATTTCTCTTAGTTGTGATAGATATAATGAGAGTGAATCAATAGAT
AAACCAATTCTTCTTTATGGCCTTTGAAAACCTGCTAGTGAGTGACTTCTTGGTCTTTTTTTCATTTTCATATTGCAAAATA
TTCATTCTAAATTGTAAATAGATGGCACCAGTGTGACTTGTAAATAAATGTTTCATTAAACATAATTTAGATATGT
AATTGTTTTGAGACTTCTTAGAGAAAAGTGCTCCCGTTGTATCTAATTCTTTCTTGTTTAAATAACATCAAGATGA
GAAATTGAGCTCTTTTTTAGTGAACCTGCTTGAGACATTCATATGGAATACAGAAAATAATAATTCCTAACATTCAGGAT
GTCATTCATTGCAATGTGGCACAGAGTTTTTTCTTTGCATCAAGGAATTTATATGACTGATACTATTTTAGTCAGGAT
AAATTACTGTAAATGTGTCCACTTTTAACCAATGGCATTCTAACATGGGATAGGCAACAGCGGGCTGTGATTTTTGCT
CTGATACTGCAACTTTGTTATAGAATAGCAAGTAACCTGATTGGCAGCTAGATCACCAGCTAGCTGGAGAAAGCAGCTTC
TAGGTCTATAGTCATAGTTGGAGTCATAACCTCATTAATGATAGCTACAATGTATTGGGCATCTACTCTTTGACAGACA
ATGAACCTTAGTACTATATACAGTTGAAGCAACTGTATGAATAAACTGTATGAATAAACTTGAAGTCGCACAGCTTATA
AAATTACAGTGCTGCAGAAAATCTTGGTCTGTGCAAACTACATCCCATATTTGTTGTTGTTGTTCTTATTGTTTTGTT
TGCTTTTCTCACTATCTAACTGCTTTGGCTACACATAGTTCTGTTTCATGGTATTAGAAAAGTAGTCAACAAGCTGC

Fig. 9.283

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TACTTATACTTTTCCTTCCTGTCCTAGACTGTTCAAGTTTCTTCTTTTAATCTTATGTATATCAAGAACATGTTTGCCT
ATGGTATTTGTCTGCCTTTCCCCCAAGATTTGTATACAAGCCTCCAACCGTGAGCCCTAGAGATATATTGAGAAATAG
AACATGGACTTTGACACGGCTCCATTCCTTGAGATCTAGCTAGTTCTGACTCGTTCAGCTTATTGACTCTCTTCCTC
TGGTCAGACTATAAATTGCAGTTTTCCTTCTCTCTGGCATTGACCAGTTAGTAGCACCATAAAACCTTTGAAATAAAA
GAGACCACAGAGAATAACCTATAATCTGATTTTCCCCACCCCTAACCCCAATGTGGAGCTGGAAACATTTTGCTAGTCT
TTAACATTTCCAGTCAAGTAGGGTCAAGAAAACTGTTAAAAACAATGTCAATGCTTAAGTGGTAAAAAGATTTGTTGA
AGAATATATATATATATATATATTTAGAGACAGGGTCTCACTGTGTTTCCCAGGCTGAGGTACAGTGGCTATTTACAGG
TGTGATCATATGCACTACAGCCTTGAACCTCTAGTCTCAATCCTCCTGCCTTAGCCTTCCCAGTAACTGGGACTACAGG
CATACACCACCATGCCTGGCTGGAGAAATATTTTAACTTCAAAAATGAAGTTTGAGCTGAAAAAGAGAGCTAAAAA
TTAGTCTGTCAGGCAAATGTGTGAAAGCAGTTCTATTATAATTTGTATTGAGCTATAATGTAATTCCTAACTTATGTC
ATATAGCTAAGTAAGTAGCTGTAATTCGCATCTTGAAATTTTAAATTTACTTCTTTCTCTCAATATAGTCTTCATCTG
AATTAAAAATTTGTGGCAATTTGTATTTATTGTGTACTTAACCTATGTGCCAGGCACCTGTGCTGTTCACTTGACCTAATTA
TCTCATTAGATCCTCATGATGAATCCTAGACAGTACTCATTAGTCTTATTCCTCAGACTGAGGCCTAATGAAGTAAAGT
GGCTTGCCTAAATTTGAGTGGCAAATAAGATAATTTTCATGATTCGCATATTTTCTTAATTATAAAATGTTACCTTTAGC
AATGATGTATATGTTAATATGTTCAATGAGTTGATCTACCTGAAGTGCTTAGCACATGGCTGGCTCCTTGTGAGTGACC
AGGAAGTGTAGCTTATATTATGCATTTTGAAGTTTACACATTGGTATGACTTTCTAAATAATCAAGAACTCTAAATG
TGAAAGCTGATTCATTTTGTAAAGGGTTTCTTTTCCATAAGGTAGTCTTTTAAACACAGTCAGAATTAAGCATTATAT
AGAGCTAGGTATTCCAGATAAATCACGTTAATAGTGTGATTCAAAGAAGACACACTGCTATCTAGGTGGGTCAAAGAAG
TAACTACTTAGTGGAGGTGGCAAATAAGAAAGTGGTCAGTGGGAAGCTAAACCACCTCTGATTTAGTTATTTAAACT
CAATGAGAACCCTCTAAGGAAGCACAACTAGGCAGGGGTTCATCTGAAAGGAATGGTCAGCTACCTGAGCATTATCTC
TAGGTGCACCTTTGTAAAGACTTTCTCCATGTATCTTCTACTGGGTGCCACCCTGTGGCTTGGTCTACGGTAATCA
CACTGCCAGGCATTCTCTACCTCTACAAATCCCTTACTCTCTAGTCCTAAGGTGCATTCCTGGTTGTGTAATTCACAG
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AGAGCAGGCATTTAGCATTCCTGTCTCTGATATCTACTATTTTATTATGAGTAGTTACCAAACAAAACAAAACAACT
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AAGTCATTGTAGCAGAAACAGGTTTTATGCATACATACAAATACATGTGTATTTTTTTGATATGTTTTGTCTATTTGTCA
CCTATGAAGAACATATAAATATTGTAAACAGAAAACTCTTTTGGATGTTGCAAATGCAAAGAAGAAAAATAAGTAAT
TCTAATTTTCAAAGGTCACCCCTGTAATCACCACCCAGATTAAAAATAGTACCAGCAGCCAAAAGTCACCTTTATGCC
CTTTAAGCCCTGCCCCCACCTACTAAAAATAATAGTGATCCTGATTTTGAAAATCACAACTTTTAAATTTCTCTGCT
TTTGAATTTTATATAAAAGGAATAATATAGCATGTACTCTTTTGTGTTTGGCTGTTTCCCTCACCCCTTGCAAATA
TGTTTGAAAGATGATCCCTTGTGATGCGGATATGGTTCATTCTCACTTAAGTATAGGTAATTAAGCAAACCTTTT
GTTTGAGATACAAATTCAGGGTAAAAATCAGATTTTCTATTTAAATAATAATTTGGTATATATGACTATTTGAGCATTT
TAACTGTTGTCAATTTTACAATAATTATTTTAGTATGATTTAATATCCAAATTCGTGTTGCTTAGATTACCCCTCAGTGG
ACATTTCTTCTGAAGGGCCTTCCTAGACTTCCCAATTAGAATTAAACTCTCTTCTCTGTGCTGTGATGAAACCTTGT
ACATTTCTCTAACTCTTGAAGGATTACAATTATTGTCTTCTGTCTCACGTACCTCTCCTGTGAGCATGTTTTGGAGTAG
ACTGGGTTTGCTCATCTTTGTAGAGCAATGAACCCCTGTTCCCAGCACCATGCCTGCTCCATCATGGGCATTCAACATC
TGTCCTTTGCATGGAAAGTAGAACAGTGAGTTCTTCCATATTACAAAGAATATTACAGTCAATTTTTTATTTTAAATTT
CTTTTAGTAGATGGCACTTTTCAGGAGATTGTACTGGGTGAGTTGTAGACACAGTAGTTACTCTTTATGGCAAGAGTTG
TATAGTATCCGGGAGGATTTTGCCTCTTCCCTGTGGCTTTCTTTTACCCTTAGTTTGCCACCACCCTGGCAATTCTA
ATCTAGGCACACCTTCCATTGCCTCCTGTGTTTCCCATTTCTAAATATATACTTCCAAGTTTTCTCTATATTTTCTGA
CTGCTCGAATGACTCTTCTCGTGTCTATAAAGTAGTAAAGAGTAGTGCATATAGAAATGCTGATAAATGTGTTGCACATG
GTTATCACCCGACTGACATGCTGACCTTCTGACTCTATCAATGGGTGATTTTTATATGAAGCCAAGAAATCTGTTGGTG
GTAAGTACTAAGACCACCACTTCAGGCTCACTCTAACCTCATTTGTGTGCATCTTGAAACCCACACAGTTGTACATAT
GAGCAGATTCACAGGAGGGATGCTGAACCAAAGTGACAAGCTTGTCTGAGCCCTAACACAGGGATTGACCCCGAC
CTTCAAAGTCAGGGAACAACCTTAGGGTTGGCTTACCTTCTGTATTGTGAACTTTTCCGATTGTGAAGCCAGCTTTTAG
CTTTACCACACTCCCAGACTTAAATATTTGGAATGGCAAGAAATTAGGCTTAAAAAAAAGAAAGGAATTAAGGCTTAA
TCCTTAATAATGAGGAAGTAATTTGTTTTAGTATTGTACAGAAACATTTTCTAGCTACCTGATCATTTAGTCTAGTTCT
ATAGCTTAAATATATAGTCTTAAAGAAAAGTTTTATTTCTGTGGGTCTATTAGTTACAAAAGTAAAGGTGCATTTTT
TATTGCTTTATTTAAGTGTTTATTAAGCTTGTCTGCCACAGGTACCCTGCTGGCTGTTAGAGAGACAATGGTGAAATA
TACAGCACTCGGCTTCATAAAATTTATGGTCTAGTGACTTTCTGAAGTATTTCTATAGGGCAGGAGACTTCTATTTTAC
CTTCTACTATATCCTTGGAAATAGTTTTTACTTTCTCTGGTAGGAGTGTGCATTTATATAAGTGTTCAGTTGCTTCCAGA
AATCCAGTGGTAAACAACATAATTTTAAACCTCCCTACTTGAATTTGGGGTGGAGTAATATAAAATAAATCTGGAATA

Fig. 9.284

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TTTGATTTTTTAGACCAAATACTTTCCCTGGGTAGAGAGATATTTTCTTTTCTTACTGGCCTGGAAATCCATGCCAT
TTTAATAAGAGATATTTTATTCTAGGTAAATGTTTACTTAAATCACCTCCTAACAAGGCTGTGTGTCAGCCAAGTTAAAT
ATTTAAATTTATTTTATTTTAAATAAATTTTAAAACCAACAGATTTCTTTCTCATTTTAATGAATTTAGCCTTTGAGTT
TTCAAAGATAAAATATACAGAGTAGAACATTTTAAATGTATCTGTAGAGAGGGGAACACAAAAAGGGCACATCAGGAAAAA
ATAGTTGGAAAAAGTGAATTTTATACAAAATTAACATCATAAAATGAATATAAGGCATTATTTCACTTATTGCCAAAAT
CTGTCTTTACAGATAATCTTCGGGAGACCTTTATTTTTATAGGCCCTAGTTAAAATATTTTTTTGAATGCTTGAGGCTCT
ATTTCACTAATAAGAGAAATAAACTTTTTCTTTCTATAAAATACTGCTACTAGCTTGTGTTGTCAGCTCTCTAAATAAAC
TCATTTTACTATTACTATGCTTGAGTGCTGTTATATATTCCTTTGTTACTGTGCATAATCCCTGGCTATAACTGCATTA
ATAAGTAAGGCCATTACATCAAATAGGGATCTAACAAAACTTTTGAATGGACTATCAAAGACCAGATTCAATGCAGGA
AATCAGAAGGAAATGGCATTATGTCTTAGTGAGTTTCAAGTTAAAAGGCCTGCCCTTAAGGATTACAACATGCCATAAAA
GGGCACCATGCAAGAATGGAAAACCTGTTTACAACAAAATATCAAATTATTATCTGACTTTTACAATGAACCACATTAT
TCTATGACTGCCAGTCATACATCTGGACTACTATACCATTTGTGTGTCAGTTACAGCAGCTCCAGTTCTATATGGCTAGT
TTTGAACCTTAGTCTGACTTTAAAACCCATCCCATTTACTATCCTCTGCTTTTTTTTTTTTTTTTTTGGTGTATATTGTGGGA
AATTCATAAATGATTTTTTATAAACTTTAGCCAGTAATCGGGAAAACTGTAAGATTTTGTGTTGTAAGCTTTATAAACG
TTACTTATAAGATTTTAATTGAATTTCTAAATTAATTTCTAATTTGAACCTCTCCCTTCTCTTAGCTTTATATGCTTGGT
CATCCCTCCTTAAGTAATTAGCTTTTTTGAAGCTTTGTTTTTTTTTGGTTTTTCACTGTTTTCTGTTATATTGACGATTTG
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ATAGTCAAACAATTGTCTAAATTTCCAATACACAAAGAAAAGTTTTAGCTAACTTACTTCAATAAGATGGATAATTTGT
TCAAAGTGCCAAACACATTTGCTTTTCGTAATTCATGCCATTTGTGTGTTAAGTGTATAAAAGAGATAGAGAAAGAATAT
AGGAAATGTAACATACACATGCAATCTTGGGCTTTCCAAAATGTTTAAAGTACCAAATTCCTTGGTGCCAATTGCTTTA
GGCATTTTCATCTTCACTCAGTTGCATCCTAAATATAGCGTAGGTGTGACATACCCTGTAAATTTCAAATTCACATAAAG
AAGTCCCAGCCTTCGAGGCAAGTCTTAATTGAATACTAAATAAAAACCTTAGTGCTCTTCTGAAATGTTTCTGAATAAAT
AGTTGCTTTACAGTCTTTGATTTCTATGTTTCGTTAGTGTCCAAATTCAGAATGGGAACCAACTTGGGGAGAAAAAGAAA
AAAAGACTCTCAGGACAATCTTTTTTAAAGGAGTAACAGATAATCTATTTTTGGCTAAGACAGGGTAGATTCAGGTAGTCA
GATAATAGAAAGGGGAGTCCCTTGTGTATCAACTGATGCTGGAATTTGTGAAATCCTATAAAGTTTTGTAGAAAAAAT
GTTAAATTCATGGAAACTTGTGTTGGTTCATAATCCTCACAGTTTCTAATTTAGTATAACATCCTGAATAAATTCATTCAG
ACCGACGGAGAGAAGAAATTATTATTTGCAATATGTGAATTTCTGACTAATAGGAGTAATAGAAAATAATTGGTCCCTT
AGGATATTTCTGATTATATATGAGAGGCTTGGGGAGGGACTCTGTGTAACCGTTGCTTTTTAACCTTGTTTTTGATGGG
TAAATAGAACATTAATGCTACATGAAAGATACTTACTCTGGCATCCTCAAGAGAAAGTAGGTTCTTTTTTTTTCTCAGTT
ATCCAAGAACTTTTAAACAATTCTAAAGAAAGAAATAAAAAGCACTTAGTTTTTCAGAAGCATGTTTCAGCTTCTTGCAT
CAGGAGTAGGCCAGATGGCCATGTTCTGTATGGCTTTATAAATATTCCTTTTTGTTCCATGTAAAAGTTAATGGCTAGT
TTTAAACCTTGGTATCATTTTCGAAGTCAGACCTTATGTGCTTAACTTCATTCAAAGTATTTCTATTTTGTAGGCTGTA
GGAAAACATAGAGCTGAGCTATTAAAATTCAGGGCAGTTTGGAAATTGCCAACAAATATGTACTGGAGAGATGGGGGTGG
ATTCTGAAGTCAAATGAATTTGGAACATCATCTTCTTTTACATGCTGGGTGATTTTGGACAAGGGGGAAAAGCTACCCAT
AGGGTTTCTGTGAATGTTAAGTGAAATAATACATGGAAAGTGCTTGAAGTAATGGTAGGGACACAGCAAACCAAAAAA
ATGCTAGCCAGTTTATTATTAGAAGGGAAAACTCCTATTAATATTTTCTTGTGTTTATGTTTGTGTTTTTCATGTTAGTCT
TAAACAAGTTACTGTTTGAGAATGTGAGAATTTTAACCATTTACAAAATGGTGTATGATATGACATGTATATGATCTC
TCACAAGTGAAATGATAATGGAAAGTTTACTGAAAATGTCTTAACAGTTCTAGGTAAAACCTTAATTTTTCTTAAATTTG
AAAATTAATAAAAGTATGAATTAGATTTAATCTAAATTTATTATTCTGTTAAAGTCACATGAATGTGGAAAAAATCAG
TATCACTTCTAACTAAATCTGGCCTTGAAACTTCTTTGACATAGTCTTCTTAAATATATATATCTTATATTCTAAATT
GTATATTATTTAGAAATATGCTATTTCTAACTGAAATGTCTGATATTTCTAACTGATATATATATCTTATATATGATAT
GATATATGATTATATATATGTTATCTATATGATATATAATATATGATATATATTTCTAACTGATATAACAATGTATCTC
AGTTCTTAGAGATATGTAAATTATATTTTAAATAGTAGCCTACAAAGATGATACAAGAAGATAAGAAATAAATATACTTT
TATATATAAAATAGTTCAACAGCAAGATTCTTATTTAGAATAATTTTCTACAGTTTTATTTAGGGTAGTTTCTATACAG
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AAGCAAAACAAATTCATAAATAAAATTTATTCCTAATTGTGTTTTAAGTCCAGTGAAGAGAACACAGGAGGGACCGAGT
TAAACAGTGTGTTTGTGTATGTCTGTGTATGGGTGTGTGTCTGTGTCTGTGTGTCTTGTGCTGTGTCTCTAGAG
ATGTACTTTAGTTGGAGTGGTCAGGGAAGTTGAGAAGTGAAGAACAGTCAAACAAACATTGGGGAGAAGGGGATTAT
AGCAGGTACAAAGGTCTTGAATATCCAGCAGATATGAGGAGCAGAAAGGCTATCTTTTTCAATCACTTAAAAAGGAAAA
GAAGTGAAAGATTGCTGTCTTAAGAACTTAAATTTATATGGCACTCATGTATAGATTTCCAGTGAAAAGTTGGACAACG
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TTCTGTTGCTAGCCTGTCTTCCAGGGGCTTGGTGAGCCGAGGTGAGAGAGAGTGGACTTCACCTAGGAAAAAAGGAA
AGGCTTCTGACATAAAAGTGCCCTTGAAAATGTTCTGTCTGTGTAATCCCTTCTCTTTACACAAAGTTTAAAGACCCTC
CCAAGAGTTTGACTTGCTAAACTACAAGAGCATTTTTTATTAGGTGTAAGATCTCACTTTGACTTTAAGTAGCAAGTGAC
ACACCAGGCAACCCTCGCTGAATTTCTCTTTGCCATAAGTGATAGATTTTCAAGTGTCTAATGCCTCTCTGAGCTAAAT
TTGCCCTCTGATCTGAGTTCTAAATGTAATTGGATAGTAGATATATGTGAGTAAAAAAAATTTCTACCAATGGTTGGAA
TAGTTTTCCCAAAGAATGTAATATTGGCCTTCAGCTGGTAATAGGATGTGCCCTGCTGGATGAATGACATGGTTGATGCA
AAAGACAGATGTAAGCCGTGGTTCCCCAACTGCAGTAGGGATGAGCATGACAGCCACACATATCAAAGGCCAGGTACT
GATATTCTTACTGGAAAGACTGCTCAGGATGTATGTGGTGTCTTTTTTCTTGGCCACCCTTAAGGAGTTTATGAATGGG
TTCCATCAGCTTCAAACAGCAAACACAGTTTGGCCTGTATGAACTGAAACCCATCACACATCATTGCCACGATCATTTT
CCTCATGTATGCAAGTTCTCAAGTATATTTTTTGGTCAAAAACAGCAAAATCATGACAAAAGCAACAATAACAACCAA

Fig. 9.285

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AGTAACTAAATACGATTTTTATTAGTATGAAATGTATTGCATTGTAATTATTTCAACATCATAACATTTATTAAAGAT
GTATGTATCACTTTATTATTCTATGCTCATCCCCCTCTTTATACAAAATGGAGCATAATCTGCACATTCATTGTTCTG
CAGTTTGTATTTTGCACCTTGACAGTATGTTGATACAAAGAGAAGCTACTTGTGTTGCTCTGGAAGAAGCTTTATTGAGAT
ATAACTCGTATGCCATACTTTTCATCTATAGAGGTTATATAATGTATTGATTTTTTAGTATATTTACGTAGTTGTTAGTT
AATATCTACCATTCTTTGACCATAATGAATAACAGGGTTTTCTCTGCACAGATATGTCCATTTAAGTAAAGATGCCAAT
GCAGCTAAATAGAATCACTCAAAAACAACTCCAAAATGAGCCAGGGAGAGTGGAGGGAAGGAAAAGGAGGAATGCATCC
AAACATAGCCCTACGTTCCATGAACACTCAGTAACATCATCAAAACGTTGATGCAATTAAATTTTACCAGGTTTACTGCT
GTCCTGATGCTTTCCAATTTTTTTTGACAACAGTTTTGCCTTTTTCAAATTCAAATGGTATAATTGGGGCTTGGTAGTT
GATGTTTATCTTAATTGAAACAGATTCTCTTCATCCTTTTGCTCTGAGACTCCCACTTTGAGGCTGAAAGGTCATTTTA
AATCTGCAGAGCACTCGAGAAGCCACATAATGAATTCAAACAAATTCATGCCTATTAAAAGGAGGCTTCTCTTTCTCAA
CTTCAGAGCTTACATAGAATAAACCATTGTGTTTTGTCTGACTTTCTTTTAAAAGGAAGTTATTGTGCTTCCTTTTT
AATGCTCTAACTACAGAGAATGGGGAAATCCTGTTTTCATTTATTCTTATCAGAAAGAAAGTGTATTTTGCTCAGTAAT
TGCAATAAAATACTACTGGGGTTACAGCTACTATACTTGTCAACAAATTTAAGGGATATTCTTTAGCTGCATTACTTTA
AGGATTTCTTGTCTATCTTTTATCTAGCAACTTCATTTTTTAAACATTTACTCTATGTACTTGTGTAAAGGTGCAAAAAA
TATATGTACAAAATGTTCTGCTGGAGTTTTTGTAAACAGAACTGGAGAAAGCCTAAATGTCAATCGGTATATGGCTG
ATTGAACAACTGATGCACAACTATAAGACAATACTGTGCATCTATGAGGAAGACCTCTCTAGTATGACACTATCTCCAG
ATGACATTTACTATTGTATTATTTTTTATACTTTGAAAAGAAGATATGCACACTCCCATGTATCCATTTTCACTAGTTTT
CTTTGTATATACTCTTAGAGTTTATGTAAGTGCAATGAAAACATAAATTCATTCTCTTCCCTTTTTTAAACAAAA
GAAGTATAATCTGCACATTCATTGTTCTGCAGTTTGTATTTTGCACCTTGACAGTGTGTCAATACATAGAGAAGCTTCTT
GTGTTGTTCTGGAAGACCTTTGTTGAGATATAACTCACATACCCTACTCTTCATCAATACAGGGTGTATAATTTCTGTA
TTTTTAGTATATTCACACACTTTTGCAACCGTCACCAAAATCAATGTAGAACATTTTACTACCACTATAAGAAACCCCT
TACCTTATAGTTATCACCCCTCCATCTCCCCCATACCTGCTCCATATCACCTGCCTCCCTTCCACCCACAGCTCTAGG
CAACGGCTAATCTACGTTTTTTTTCTCTATAATTTGCCTACTCTGAATGTTTCATATAAATGGAATTATATAATATGAG
GTCTTTTGTGACTTGCTTCTTTCACTTAGCATAATGTTTTCAAGTCTCATTTCATGTTGTAGCATATATCAGATTTCAAT
TTTTAGGACCGAATAATATTCATTATATGAATATACCACATTTTGCTTATTTATTTCATCAGTTGATAGACACTTGCCT
TGTTTCACTTTTTTGGCTATTTTGAAATGTGCTGCAATGAACATTTATGCAAAAGATTTTGTGTGTGCATATATTTCTCT
TTCTTCTGGGTATATTCTCTAGTAGTGAATTGCTGGGACAAATGGTAGCTCTAAGTTTAAACCTTTTGGTTATTTTCCAG
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TATTTTTTAAGTGGAAAAGTTAAGGCACAATGTTCCAGTTATGGAAAATATGCTAAGACTGTACATACATAAATATGT
TTCCTCAAAAGCATCTTTCTGCGTATGCACCTAAGGTGATGGTAGAAGTTGCTAGCTTTGAGGAAGAGGTGGCTACAC
GTTTGGAAATCTGAGGTAGAAGGGGGCTGAGTACTCACCATTTTAAACTTGGGGAAATTTTACTATGTGTGCTAATTAAT
TTTTCAATTGAAAATTTGATTCAATTAATGTAAAGAAGAAGAAATGTAAAAAATAAAAAAGAAGAAGAAGAAATGTA
TTTTCTGACTCCTCTGACCAATGTTGGTCAGGCCAGGGGCTTCTTGACCTGTGCTTTGTTTGTAGTCATCCGGCACCA
TGAGGTGCTTAGTGCTGTACAAGCACTCTTCTTGCCATCTGCTCTGCTTGACATCATTTTTAGTTATTAGGACAAAAAA
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GGGATTGAGAAATAGGAGGGTGGTATGTGAACATAATCAATTAATGGTATTAGCAAATAGAGCTGGGCAGTCTGTCTGAAG
TGCTTCCAGTTTTTTGTACATTCTTCTCCAAGTATCATGTGTTGGTGACTGGAAGAGGGAGTCTTACTTTCCGCACA
GTTTCCATCATGTAGCTCTGGAAGGGCTTTGTGTTTGTCTCATCTTTGCTTGAGAATCAGCTGTCTTATTTTGGGCTT
TTCTATTTTTTAAGACAAAACCTACCTATAGTGGTTATACATGAAACACACCAGTAGATTTGATTTCTGCCTTTGGTTA
AATTAGTTTGCCTTATTAAATGATAGGAAAGAATCAAAATTCGTTTGCCTTATAGATTTGTTCATTCTTATATTCATC
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GGGTACCTAGGAAAGAAAGCAATAATGAAAGTATCACTTGGGTATTTGTTTTATTCTGGATCTTATTCTATCTGCTCCA
TATTATCACTTTGGCATTTAATAATCTAGATGATGGCTGGGTGGGTGCAGTAGCTCACACCTATAATCCTAACACTCTGG
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TTGCTTGAGACTGGGAGGTCAAGGCTGCAGTGAGCTGTGATTGTGCCACTACACTCCAGCCTGGGTGACAGCAAGACCC
CATCTCGGGGAAAAAACCAGATGATATGGTTAACCATATTCAGTCATTATTAGTTATTAGAAAATAAGATTTA
CAAGGCATCCATGGAGGGGAAAACAATTACACACCTGGTTAGTTGGGCTGAGGCTTGACAGAGATAAAATCACCTGCACA
CTGTTGCAGAGCCAGTGCTCCAACCTAGACTTCCGGACACCAAGCCTATGGCCATTAAGCACTCTGCTGGACTGTATC

Fig. 9.286

TTGGATAGTTTGGCTTTATGGGGAACGTAGTACAACCTTTACAATACAACCTTTAAAAATAAAGTATAGCAGAGTAGCAGTT
TGTCCACAGTCAAATATGAAATATGTAAACATTTACAGGTTCTTTTTTTAATTTTTTATTTTAGGTTTGGGGGTACATG
TGAAGGTTTGTATATGGGTAGACTCATGTACAGGGGTTGGTTGTACATATTATTTTCATCACCCGGGTATTAAGCCCA
AAGCCTAATAGTTATCTTTTCTGCTCCTCTCCTTCTTCCCACCTCCCTGATCAAGTAGACCCCAATGTCTGTTGTTTC
CTTCTTTGTGTGCCTGAGTTCTCATCATTTAGCTCCCACTTATAAGTGAGAACATGCAGTATTTGGTTTTCTGTTTCTG
TGTTAATTTGCTAAGGATAATAGCCTCCAGGTCTATCCATGTTAAAAGACATGATCTCATTCTTTTTTTATGGCTGCATG
GTATTCCTTGGTGTAAATTTACCTCATTCTCTTTGTCTAATCTGTGACTGATGGGTATCTAGGTTGATTCCATGTCGTT
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TGCCAGTAATAGGATTGCATGGTCAAACGGTAGTTCTGCTCTTAGCTCTTTGAGGAATTGCCACACTGCTTTCCACAA
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GCTGTTATGGTTACTGTATCCTTGTATTATAGTTTGAAGTTGGGCAAGGTGATGCCTCCAGCTTTGTTATTTTTTTATTA
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CAGGTGAGGTTCAGTGATTAATAACACTGGAAAGGAGAACAGGTTCAGAAACTATGGCTCAGGCTACCTATTACGCC
ATTTTCATTTTTTTGTAATGCAAGTGCCATTACATAATCAAGGGTATCCTTAGTAACATATGAAGCCTACATTCTATTT
CCATTTTTTAAAAAGTTCACCAGTAAACAATTGTACAGCAAATTTTATCAATGTAAAAAGCCATTGTACTCTATCCAGTC

Fig. 9.287

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AAACCATATTACCTCTCCTAAAGTGCCCATACGGAGGCTCTGGGAAATTGAAGTTGCCCTTAATCTTGAGTTACAAT
GGGCAGGGCCTCTTTTTTCTCTAAATTTTACTCAATAAATGCAGGCTTCCTATGCATTAAATGGTGCCCAACAAACATT
GAAACTACTAGCTCACCTCCTGAAATTCAGCACTTTACTATGTGTCTTTCAATGTAAGAGCATTCACTAATTTAACAAG
CATTACATAACATGTGTCTTAATGAGTTTCACTTAGCTAGGCCATGGAATAGATATTCCTGTAAATCAACTCCTTTACA
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TCTGACAAAGTAGTAAAATAGTCTCACTTCATCTGGATAAAGATCACCATCTGGAACCTATAAAAATTGCAATGATTCCA
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TATACATATATCTATACAT
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TGAAACCTACGTCTCCTGGGTTCAAGTGATTCTTGTACCTCAGCCTCCTGAGTAGCTGGGATTTTAGTTTTGCCCCACC
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GATCAATTCAGAAGTTGAGTCTCTGTGAAATGATTTCAATAGTTGAGGTGATTTTACTGTTTCTCTTTAATGTTGTG
ATATATTTTCTCTCTTATACGACTCTATAGTAAAAACGAGAATCATTTTACTCAATCTGGTTTCAATGTAGCAGTATCAGG
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TGTAAGTCTGATCTTCATCCCTAAGACCTGATTCATTTGCTTTAGATAACACTGAGAGCTAATCATTTTTTAGCAAG
CATCCCAGGTAAATCTAAGGCCATATTGTGAGAAAAGCCAGTATAACGATGGAGAAATCTTATGTTGATGCTCTGACAC
TGGCTCTACATCTGTCCATAATTTATTTAACTCCTCTCTGCCTCAGTTTCTTATCTATAAAGGAGGAAAGGAAATGCC
AGTCTCTTCCACCTGGGATTCTTGTGAGATTAAATGAAATAAGCCATGCAAATGATTTAACACAGTCTACAGCACACAG

Fig. 9.288

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TAAATACTCAATAAATGTGAACTCATTATCGTTACTGTTGTCATTGGTATTCATATTGATATCATTATTCCTGCATTGG
AAAATGTAAATGTACAATTTTAAGTGATTAGCATTGCTGGTGGATTATTGATTGCAATTTGTCCTGCTGTCTCTTCTGA
CATTGCTAAATTTCAATTTACTTAAATATTTCCCTTATTTTTCTCTTGTTCACCCCATGCACCCAGTGGAGTTGAAAT
TGAGTTGAGATCAAATGCAGCAGGTGTTGCTCAGAGAATTTGGTAAGACTAGTTGAAAAAAGATCAGTGAAACTTTATC
AAAAATAGAATAGTGATTCTCCTGGTCACCTGCTTAGAGAACCCATTAAGAAGTGTGAGGTTCTCCAGGCCACCATAGA
GCTATAATCTGCACCTTGTATCAGCCATAGCAGGTATTTGCACAGTAAATTTCCCTCACCTAGTTTATTCATAGGTCT
GATCATAGCACACTACAGACTCAAACCTCCTGGGCTCAAGTGATCCTCCAATGTCAGCTTCTTGAGTAGCTGGGACTACA
TGTGTGGGCCACCATGCCTGACCATGGAGTTCTTAAAAATGGATATTGACATACATATATAGAAAATTCATTCAATAAA
TAGTTATCACTTTCTTATACCTGGTGTGAGAATTATGCTAGATATTGGAAATACAGAAATGAATATATTAATATTAAAG
TCACTTAACATAAGCTTTTCCCTGTGATAATCTTTCTGAAACAAAGCAAATGATACAGAATTCTTTAAGTACTATTCAA
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GATTTGGCTAAAGAATCTACTTTGACCAAATAGGATACCTTAAATAACAACATCAGCAAATATGTGTAAAAATCCCCA
GAGAAAAGCCCTCTTAAATGCCTACTTATTTCAAGTCAAATTTAGTTAACTACATAACTTAGTCCCTGAGTGTGCTAGT
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CATTTCTAAAGATGGCACTCTTTCTCCTTCCAGTTCTTTCTGTTCAATAGTAGGGCTGGGGAAAGGCAGAGGCAGTTT
TTGTTATTTCAAATGACAGCATCAAAGATAGTAATCCACGGTGCTCAACAAAAGTCGAATGACCTTTTTTCTCTCTCCA
TTCATAAATACATAAGAGCTGATGCTTCATTATGTTTAAATACAAAATGCACACTCCTACTTTGTTTTCTTATATGTGA
GTTCTCATGTATTCTTCAAATGCTCATCTAATTACTGTTACCTACTATTCCAAATGCAAATGACAAGGTCCCAGTTTAC
TGTTGATCCTATATTACAAGAGTCATCAATTTTGGTTGAGAAACACAAAGGACAAATATCTCATTATTGTGCTAACCAT
GCCTATTATTAGTTTTTGTGCCCCCATACATAAGTAATAGCCCCAAATACATGGCAGTTATCACACACCAGGCATAATT
CTAAATGGCATGTTCTCCATGTTATGGAAGAAGAGACTGAGGCATGGAGAGAGTGAGTAACTTGGCCAAACTCACAAAGG
CTACAAAGGCAGAGAATCAGGTTTTGAACTCAGGGCTGCCTAGACCCTGTGTTCTTAACTATCATCATATAGTGTCTCT
CTTACTGTTTCTCACTGAAGATGAGAGAGGATTAAATCTCAGGAATAAATGTAACCTCCACAGGTAGGCTTATGCATA
AAATTCAGATGCAATGAATAACAAAATGACTGCCTCCACAAAATTAAGAAGCAAACATATAATGAGGACTTACTCTGT

Fig. 9.289

Fig. 9.290

ACCTCTCTTCATTCCCCCTCCTCCTGGTAGCTATCATTCTATTACCTTCATGATATCAACTTTTTTTTAGCACCCACTTAT
GAGTGAGAACATGCCATATTTGTGTTTCTATGCCTGGCTTATTTCACTTAACATTTCAGGTCAATTCATGTTGCTGCAAA
TGGCAGGATTTTGCTCTTTTTATGACTGCATAGTATTCCATTGTATGTGTATGTGTGTGTATATATATATATATATG
TATTTTCTCTATCGGTTTGTGTTGGTGGATACTTTGGTTGATTCCATATCTTAGCTATTGTAGATAGTACTGCAATA
AACATAGGGGTTTCAGGTATCCCTTTGATATACAGATTTCCCTTTCCTTTTCCCTTTAGATAAACACCCAGTAGGGGGATTGC
TGGATTGTATAATAGTTCTACTTTTAGTTTTCTGGGAAATCTCTATGCTGTTTTCCATAATGGCCATATTAATTTATAT
TCCCACCAACTAAGAATTCCCTTTTCTCCACTTCCTCACCAGCATCTATTTTTTTTTTTTTTGGTCTTTTAATAATAGC
CATTCTAAGGTAAGATACCTCACTGTGGTTTTGATTTATAGTTCCCTGATGATTAGTAATGTTGAGTACTTTTTACATA
CCTGTTGGTCATTTTACATCTTCTTTTGAGAAATGTCTTCATGTCCCTTTGTCCACTTTTTTGATGGGATTATTTGGGTTT
GGTTTGGTTTTGTTTTTGTCTGTTAACTTGAGTTCTTGGCATATTCTGGATATTAGTCCGTTGTCAGATATATATTTTGC
AAATATTTTCTCCCATTTAACATGTTGTCTTTTCACTCTGTTGATTATTTTCTTTGCTATGTAGAGATTTTGGTTTTA
ATATAGTCCCATTTGCCTGTTTTTGTGTTTTGTTGCAATGCTTTGACGTCTTAAGGGAAGCCTATTTTCGATCTGGTACTT
CTGAAATCTCATTGGCTCAAGTGAAGCACGGGTGTAAATCCTGAGTGTGTTGTTAATCTCGGTTCTGTGTGTCTTCTGTA
ATCCAGAGACAAATTCATGTGTGTCAGGTTACACTTGAGCTGATCCTCTGTAACCTTTGCTGATTGTTGTTAAGACAGAAC
AGAACCAGACTGGCTTTCACTTTACACAACAATCACAACCTGTGTGTAATGATTATAGACTTATTCTCTAACATCTCTTA
AGCACTTAGGTAATGAGAAATGTCATAATGAAAATATCTGTTAACCAGGGAAGATCAAGGGAACAAAACAAAAACCTA
AATGAAGTGTGAAGGTGGTAGAAGTAAAGACATAGATGGTTTCACTTGGGCCAGAGATAGATTGAATGTGAGACATTTG
AAAGTCTAATGTACTTTCAGTACGTACTTTCCAGTTTACTTTGATGGAACGCACTGTGTTGAATGTTATTGTGCTGACA
AATAAGCCCCCAAATTAATGACTTAAAAATAAACTCATTCTTCATCAAGTAGCGTTCTAGGATGGTGTGTTAGGAGGA
TGTGGCAGCTCTCCTTGCTCTCAGTCACGGACCAGGATTCTCCCACCTTCTGTTTCTACCATCCTTCACGCATCAGCAT
CTTTAGCATCCAGCTGGCAGAAAAAGAACATCAAGATAGGCATGTGGGAGGAGCTATGAGCCAGTCCTGAAAGTGGCAT
GCAATATTTCTGCTCACATCTAGTCACTTGGCCAAATCTAATTGCAAGGGAACGAGAAGCTTCTTTTATGCCCAGGAACA
GGCAGAGAATGGAATATGGCAGACAACAAGCAGAATTTCTTATACTCACAGTCCTGATTATTGACTTCTATGCTTTCCC
AAGGTCATTTTGTCTTCAGCCAGATTCCATTTCAAGGAAACATGAAAATGTTTCTTCACTCTATAAAATCATTGTTGAA
GTAGATCCTGCTACCTCTGTGGTAGCCCATCCAACATCTTAAACTTTTAAAGATAAATTTGTAACCATAACATAATTCA
CCCAATATAAGAAATCATGAGAGAAAGCTAATCTCAGTGTACTTATTATACAATGAGACACATAATGAGACATATCTTT
TTTTATTCTCCACTATATATTTTAAAGAAATTGAAGAGGCAAGTGATTGTTTATGGCCATCGTAAGATAATATTCTTAT
CATTGCTTTGCCCAGTAAGATTGAAAAAATTGTCTGTAATTCAGGCCACACAACCCTCTGAAGCCCTTCGCCATTCATT
CAAGCCCCATTAAGGCAGAAAACCCATCTGTTTGGATTGTTGAGGTTGGAACTGAATAATATCACTTCTCCAAATAG
ATTTAATAGTAGGGCTGGTGAATGGTTTCCTGACCTGTTTGTATGCAGAGTGCAAACCCAGAGGAAAAACATGGTATATG
AGTTTTCTGTATCCATTAGTCTAAAAGAATCAGAATTCGGTTATATTTTAAAGGCAATTATAGTAGAACTATTACTTTTT
TGTTTTCTGTTATAATCTTACCTAGACTTATTTAACATACTTAACACATAATATTTTAAAGGCTAAGTATTTCTACAAAG
GATTTATCATTCAATCATTCTTATTAGTCAATAAAATCTTTTGGAGATTAATTTTATAGTAAGCCTAAATACCCAAATA
GCCAGGAATGTGATTGAGGATCACATTTTAAACCCATCCCTCAAAAAGAAAATTGTAATATCTTGAGAGACAGGTAT
GGTTTGAAGATCACCCCTCTTCAAAGTGAGTTCAATATCTGACCTAATGGAATCACTCCCCATTCCCCAGGACTAGGTG
ACACTCACTGATCTGGGAAAAATAAACACGTGCAACTAACAAGAGAAATTTTGAGAATTATGAGTAAGCTTTGAAAATT
TGGGTACTGAAAATAGAGAGGGGAGGGAGAGAGGGAGGTGATAGGGAGGGAGGGAAAGAATAAAAAGGAATGAAGAAAA
ACTAATAACAATTTATCTCTTAAGAAAATAGAAGGTGCAGCTGGGTACGGTGGCTCACGCCTATAATCCCAGCACTTTG
GGAGGCCGAGGCAGGCGGATCGCCTGAGGTGCGGAGTTGAGACCAGCCTGACCAACATGGAGAAACCCCAAATACAAA
ATTAGCCAGGCATGGTGGCGCCTGCCTGTAATCCAGCTACTCGGGGGCCTGAGGCAGGAGAATCACTTGAACGCGGGA
GGCGGAGGTTGCAGTGAGCCGAGATCGCACCATTGCACTCCAGCCTGCACAACAAGAGCAAACTCCATCTCAGAAAAC
ACATAATTAATTAAGTAATGATATAAGGTGCTAAATTTTTATTTTTCACCCATCCAGTTTCTTTTCTTATTCTTTCTGA
TATGTAATTACCTCAGATGCAGATCTGAGGTGAACTAATGAAGATCAAGCTAAGAGCTTCTCACTGGCCTGGTTCCCT
TTCAAGTTGTAAGAAGTGGTACTAGCAGCTGCACGTAGTTTTAGGTTTTGTAAAATTCAAAAACTAAGATTTTTTTGT
ATTATTTTTCTGAAAGCAGACCCTTATAATTGTATAATCTTCGTGTACCACAAAACCTTGATCCCACCCCTGATTGCAT
GGCTGACTGCTGTTCAAACAGAAGGATATTCAAATAAACCCCGTTAAATGCCTTCTTAGAGATGTTCCAGATTATTT
CTTCAAATGTGCTAATCAATCTCATTAACTATTTCTTTTAAATAAGTGACCAACTCCTAGCTAAATTAAAAAATAGTTA
TGAAGTTTATTTAAAGTAGAACTACACAGATAACCATGGTAAATGATAACCGGTATAGAAAAAGTACCGCTGCGTCTAA
AGATACCCATGTATTCACGATACAAATATTTATTGAGCAACTCGTACGTGTGAGGCACTGTGTACCTGCTGGGGGACA
CATTAACGAACAAAGTAGATTTTTTAAAAAAAATCTCTGCACTTGTGGAGCTTATATTCTAATGGGGTGAGTAAGATGA
TAAAATAAGTAAAAAACAAGTTTCATCAGAAGCTGTTAAATGCGATGGAGAAAAATTAAGAGTAAGGAAGGTTTTTTGT
TTGTTTGTTCAAAAAAGGTTAGGAGGTTAGGGGTGCCAGTTTTTAAATAGGCCAGTATGGGAATATCTCATGAA
GGTAACATTGAGCTGATATCTCTAGAAAGAATATGCCAGGCAAAGAGAACTGCAAATCCAAAGGCCCTGGGGAGGGAGT
ATGCACAGTGCTTCAAGGACAGCTAGGGACCAAGAGGCTGAGCAGAGTGAACAAGAGGGAGAGATGTAGGTAAAGATGA
GATTAGAGAGGCACCAGGGACCAGGTCACTGAGGGCCTTAAACCATTTGGGAGAACGTAGTATTTTCTCTGAATAAAATA
GGACACCATTATGGGATAATGAGTAGGAACCTTGCCGAGAATGGGCTTACATTTTAGGAGGATCGCTCTGTCTGCAATAT
AAAGAATAAATGTGGGTGGGTGGCTGGAGAAGTAGCAGGGCACATCATTAAATATCAATGTTTCAAAGTCGGCTTATT
AAATCAATTTTACTTGCTCTATGTCATAGAAACAATAAATAAAGACAGAAACACCCCGCAACACTACCTTGGGTATTTT
TCTTTGCTTGCTTGTTTTTTTTTTTTTGTGTTGTTGTTGTTAATATTTTAACTTTTATTTTTAGGTTTAGGGTTACATGTGCA
GGTTTGTATATAGGTAAACTTGTGACCCCGGGGGTTTGGTATACATATTATTTTGTACCCAGGTGCTAAGCATAGTA
CCTGACAGTTAGTATTTTTTTTTTCTGATCCTCTCTCCTCCCACCCCTCCATCCTCAAGTTGGCCCCAGTGTCTATTGT

Fig. 9.291

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ACCTTCTATAATTACTAGGTTTTTGAATGACAGTTGAGTGGTGC GTTGTATTCAGGTAATAGCATTAGGAGCCCCCTGGG
TTAAGGAGTGAATTGAAAGTGGAAATGTGAGCAAATGCAGGCATCTCTTTAAAAGGCTGATGGGTGGAGTAGGAAGAAA
GAACTAAGGCTGCCTGACAGAGAATTAGGTTCAAGCGGAATTTTTTTGAATTCAGGTTACTTGAGCTGAGGGAAGGCTC
CAGGAGATAGAAAGAGACTAAAAATTCAAGGGGAAAGAGGGTTTTAAAGATGGAGCTGTAGTTGCTGACAAAAACATGAT
CATGAGCCAAAGTAAGGAGGCTATCCTTGGAAAAGTAGTAAATACCAGAACTGTATTCTTTAGGAAGGACAAAGAAGAC
CTGCCTTGGCAACATAGTGAGACCTTGTCTTACTAAAAATCCAAAAAATAGCTGGGCATGGTGGCACATGCCTTTAG
TCCCAGCTACTCTGGAGAGGTAGGAGGATTGGTTGAGCCTGGGAGTTTCAGGGTTGCAGTGAGCTACGATCATGCCACTG
CAGTCCAGCCTGGGTGACAGAGGAAGACAAGACTGTGTCTGGAAAAAAGAAAAAAGAAGAAGGAGGAGAAGGAGAAG
AATAAGAGAGAAGAGAAGAAGGAGAAGGAGAAGAAGAGTAAAGAAGGGAAATGTAAATTTAGAGATGGAGGAGTAAAA
TGTTAAGAGATATTAGCCATGAGTTCTCTAACATTTTACTCCTCCATCTCCAGATGAATGAATGTAAAGTAGAAGGT
ACAGCCATCAATAGCCAGAGGAGAGAGGGATGGGGCAGCTTGAGAAGAAAGGGAAAGGCTTAAAAAGCCACTATGCAG
ATCAAAAAAGGGAACAGGGTAAAGGTGAGTAGAATACTGACCAGCCCCATAGATAACAATAAACAATGTAAATAGGCG
AATGACAGAATTGAAAGTCATCTAATGCAACTTCATCAAAGGTGAGTCAGGCTTGGTATTGACAAAAGAAAGAGGAAAA
CTCACAGTGAGTTAGTGGAGTCCATTTATGTAGTTATGTGTTCTACCTTTTTAAATTGTAGTAAACTGAGTTTGGGATA
GATTGTTTCTTTCATACATTCTACTCCAGTTAGTAAATATTAAATATATACATATATTTTATGAAAAGCTTATAGCATT
TCATATTTAAATATGAAATGCTTTTATTTCAAATCAAACCTTGCAAGGATACCTCATTTTGCTGTGTGCCTCAAGAGTT
CTTTATCTGATCAGACTCAGGTAGGAATGATGAGTTTAAATCAGACCATGAGTCAACACTATATTTTGCTGAAAAGTAAT
GTGTWGATATGTTTAGCTTTCCATTCTCCTTGGATATTTACATTGGAGCAGTAAGATATTCCTTTGATACCAACTCCTA
GAAAGCATTCCTTCTACCAAGGAGATTAAGTTTCTGTGGAAGGAAAAGGAAATAGAATAAATAAATAACTATTTCAACAT
TTAATCTGACACGACACATATCTTCCACCATAAAGGACTGTGCCTTCTCTTAATTTGAATTCATATGGCCATGTGTA
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CATTAACCAATGAAATACATAGCAGTATCAAAATTTGATTAAGAATAAAATAAAAGCTCAAGGAGGTACAAAGTAAAT
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TAATTCCTAAGAAAAGTAAGAAAAAGAAAAATGTAAAGTATTAGCATGAGTCTAGTAAATAAATACATGAAAATAAACT
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CATGAAGTAACATGAAGAAAATTACAGCGTAAATAGACAATATTTTTGTCATCTTAATATTTTCAGCATCTGAATAACAA
TCACTTCTGTGTTTTTCACATCATCCAGTAATATAGTATCTGTATTTGGAAACACATAAAGTTATGATACATTGCAAAGA
TTAAGAGGAGAATTAGCAATGACAAGATAAGTAGAATTAAGTCACAAAACCTATGAATTTATTAAGTATAAGTTACCTGT
CAAACCTACTTAAGTTTTAAGAAATACTCAATTCACCTGGATAACTAATCCTCTATTCCCCATCCTCTGTAAATTAACA
CCAACACATAGAATGCAAAGATGATCCAATCGGCTCTTCACTCACATTCATATACATTGCTTCTACACACAGAATATGG
GGTGGGGTGAATGCATTTCAGAGGGCCAGTGACGAGGAAGTGAGTAGATTTGATATACCATAAAAAGCTGGGGCATATTT
GGGAATCTAGCTAAGTTATCCCCAGGCACCTATACAGATATCTCAGACCTCACTGATCAGGCCTTCGGGGTGCCCCATTG
GGGCATTTTCTCAGCTAGTAGATAATACTCTCCCTACCTCCCATGTCAAACCTACTGTCCCTTCTTTAGCAATTAACAAG
TCCCTCACCACCTTCTTGGGCCTCTAATGTGTGTTTTTCAGTCACAGCCACTCCTCTCTACTACTTTTGCCTGCTAAGTAT
AAGAGTAGTATTGCTTCCAGTCCTGAGTTTGAAAAAGGAAGAAGAATTCCCTTTATGTCAATTTGTATCTTTTATTCCA
GATTATTAATCTTCCCTGGTCTCACTAGGAAGAATATAAATAAAAAGACACGTTGCATTACAAAGGTTCTAATTCTCAC
TGGATCTACATGAAAAGTTTGTAGTCATCTCTTACCTAGATTGCAAGATAATATTACACTAATGAACATTTCAAAGCA
TATCAATTTGTTATCTTGAAAGCTTTCTTCCATTAACCACAGAAAGAAAGCTACCATGGCCAAATCATCTCTTCCCCAG
TTCTGGGTTTTTTCCCAGCTGGGGAGGTATCCTCACCCCTCACCCCTGTGGTGTAGGAGTGATGGGACTAGGGCCAGAGTT
CTCAAAATGTGTCCTGAGGCAACAGAGGGAATGTACATGGGTGCAAGCGTTATTTTAAATGTTCAAGAAAAACACGAAA
ACATCTGTCAACCACCTGGAAAACCTAATAGCTTGGAATAGTTCCAGTTTCCAATTAAGACCATGATATTCCTTTCTGT
GACACCATATCTTTGCAAAATTGGGTTTTTCAGCAGTTGCTGGATAAAAATGAAGTGCTGAGCAGAAATCAATGTGCAAC
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AACAAAACCTTATTAAGTTATTTGGACTTAACCTATTTAATAATTGGAACCTATTATTTATTTTGGCCAATGTGTGCTGT
TATTAAAAAATTGAGATATTAAGGGCACCATGTACTGAGAAAGTCTGAAAATATTTGGATTAGTTGTTTATTCATATCT
GAAGAATTAGAGCTAAATCAGGCTAGGCATGGTGGCTCACATCTGTAATCCTGGCAATTTGGGAGGCCAAGGCAGGCAG
ATGGCTTGAGCTGAGGAGTTTGAGACCAACTTGGGCAACATGGTGAAACCTGTCTCTACAAAAAATACAAAAATTTGC
TGGGCATGGTGGTGGGCGCTGTAGTCTCAGCTACTTGGGAGGCTGAAGCAGGAGGATCACTTGAACCCAGGATGTTGA
GGTTGTAGTGAGCCACGATTGAGCCACTGCACTCCAGCCTGGATGACAGAGTGAGACCCTGCCTCAAGACAAAAA
TGATTAAAGATAAATCAGATCTCTATGTGAGACTATTTGAGAGAATTTTGTAGGAGTTAAAAAATAACAGTTTCCAA
GCTGTGAAGTATCTGATGTGCTCTAGAGTAGAGAATTTATCTTTTTCTTAATTCGTGTTAGGATCCCAGGTAGCCAAAT
CTTTTTTTTATAAAATGTGATATTATTTTGGTAATGACCTAGATAGACCAGGCTTTTTCTTCCATTGCTGTGTGCTGGCC

Fig. 9.295

Fig. 9.297

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TCAAGGCCAGTCTGGGCAACATAGCAAACCTCCATGCCATTGAAAAAAAAAAAAAAAAAGGATTAGATTCAAACGTAAGCC
CCAGCCTGTGACTCCTTCACAGTGCTAGTTCATGCATTCATTTGTCATGCTAAAATTTGCAATGTGTGATATTCAAAG
CAGTGGGTCTTACATGGCCACTAATGCTTTAAATATAGTAGGATTAGGGGTCTTTGCTCCATAAAGTAAAACTTTCTT
CTAATTAACGTATCTCCTTTTTCCTTATGATTTCTGTAATAGTAAAACAGATAAAGATTGCCACCTAGTGTGTTTAAGAT
TTCAACCAATCATGAAATGAACTTCTATAGGAAATTATATTTTCAGGTGCTGATGTTGGTTGAACTTATTTTTTTAAGA
TGACAACAGGCTTTTTTCATTTTTTAAGTACTTGAACCATATATGGATCTCAACAAAGTCACCATAAGAGTGTGGGCATTT
GGATTTCTTGCTTTTAGTTCCTTGCTTGTTTCTGTTTCCTTATTACAGAAAGAAAAAGGGTAGAAAATAACGTGATAAA
TAATACAGAAGTAGAAGAAGGATGGGAGGGGGAATAGAGAATAGAAGTTCTTGACTCTACTGTCATTTCTCTTGAGAGA
AAAAAAAAGACAATAGTATACTGTTGAGCCCTGTATTCTCCTGCTTTCAACAGCGCTGGCTCAGGACTCAGCAAGTCTT
TCTCTTTGGGAGAGACAAGGCTCGCCAGGACTAGTGAAGATGACTCTGGTTCACCTCCTGGTAACCAGGGAAGGTGACA
GAGCATGTGGCACCTCCTTGACATCATGAAATTTAAAACCAATGTGACACTGACTCCCACAGTGACCTACAGTGGCTC
ATCCTGGCTGTTCTAGAAGTTTCTGCACTTGCACTTCTGCTGTGACTGTTACCTTGGTACCAATCTCATCTCTCACATAC
ATACATATGTGCATACATACATACATGTATACATGTGTGCATACATACATACATATAGACACGCAGTGTCAAATGACT
TTTTTTCTGAGCCTTCCCTATCCCTCATTCATTCTCCGCTGTCTCCATGAACACTCCAGTTCTTTTTTTTAAAGAGC
TGTATTTTCTTAAACAGATCACAGCTGTTTTTTGTTAAGGGTGTGTTGTTGTTATAATTGTTTTTATTTAAATGAAAG
ACAGTATTGCATCATCCTCACCCAAAATTCACATTACCAGGCGGTACAATTTCAAATGTCCAGGCGATATTCAGTGTG
TGACTTTGAATCTAGCTTTTCAGTCACTTTATAGAGACCCTAATAGTCTCTTCCCATCTCAGAAGAGACTCAGAAAAAT
ATTTAGCCAAGTAGGCCTAGGAAGCCCCAATTTATTACATTGTATTGTAATCCCTCCCAGCCCCACTTTACGTGGCAAG
GCGCTTTAAAAAGCTTCTGTGGGTAAAGTTGTTGAGGTGCAAGATTTGAGAATTATTTTATTTTATAGGATTTTTTTGA
GCATATGTGATTATTAGTTCATTAAAACCCACACATACTTTAAACCAAATCATATTCACAAAGTGAATCTTCTGTTGA
TCTTAACCTACCAGATCTGTTTTTAAATGATTATAAGCAAAGTGTAAGACACAATTCAAAGATATGTTTGTATTATATTT
ATTTTCTTCTAACCACAAATAAAAATAGGTTTTACTTGCTTATTCTCCTAGTGACAGTATGTCTGAAAAGATCTGTTT
TAATATTGAGTTGAGGACTTAGACAAAGATTTTAAATTTATATGTTAGGTATATATTCAAAGATAATTTAAGCATGACA
TACTTTTGAAAATAAATGGATAGGCTATAACCTGCCGTTTATCGAGGACATCTGCCACTGAAATGCTAGGTATCATTTT
ACTGTTACTAAAATGAGACCATCTTGAGATATTTGGCAGAAACAATTTCTGGGCAATTTATTAAAAAGTAATATATTTCT
CTTCACTAAAATGATAACTGTCTTTATGCTTTTCATAGGGAAAAAAGGCTGGTAAATTATCATAAATATTGTAAAGTGTA
TTACTTATAATACTTATAAAGGAGATACAACCAAAGAGAAAATTAATGTTTTAATAAATATGCAACCCTTTTGAGTTT
TGAAAGATTCAAATCCACTGCAATCTATATAAAAAATTAGGAGACATATTTCCATGGATTCAATTCAGATTGATGATGT
GCCTTGTCAAGATGACTTAGTCTGAAAAAAATAAACTTTATAGTAGGTCAATTATAAAGAATAAAATGAAAGAAGAGA
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CAATGTCTTCAGCTTGGCCATGCACTGAAGCAGATGAATAGTTTATCAGACTGGCAAATCAAGTCGGCATAGGTGGCAT
GTCTTTTCTCCTTGTGTAATATACCTCCCATTTATGGAATTTTAAATAACATTTTAAATGTCAGAAACCATTGAACTAT
CTTGAATGCAATATATGTAGTATATACACCTATTGAAAATGTTTTTTTGTCTTCCGACTTCTAACACACTACACAGTTTA
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AATGCTGAAGTGTTTAGCTCTGGATTTTGTCTTCATGCTTTTTTAAACACTGGTGTATCTGFAAATATTTCTCAGTTGTT
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GAAGGGAAAGTCAGAATGATGGGCCAAGGAAGAAATGTAGCTGGCCAACCCTGGCATCACTTTGTAATTTGCATTTATT
GGGCCATTTTCTAGAGCTTTATTTTTTAATCTTTTTTAAGCTGTTCCATAGGAATCAGCCAATGCATATCACAAAAGGGG
ACCATTCCAGGTATGGCTGTGCAGCTTACTACCTGCGTGTCTTTGTGTTAAATCTCTTAATCAATTTGAGCTACAATTT
TCTCATCTGAAAAATGGGAGTAATGTCACCTACCTTGCAACATTTGTTGTAACAGCTGATGATTGTATATAAAATACACT
TGCTGTGATAACATTTCTTTTTACTTTTTGTAAATGCTTGAAATTTTCATATCTTTTTTTTCCCAGAACTTTTTCTTTCCC
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TTTTTCAGCACTTAACATTATTAATTTATTGAATCCCAAAACATCACTCTATGTATGTGCTGTTATCTCCATTTTACAGA
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CCCAAACCTTCATCAGAGAGTAAAATCCAAGTTCTTGTACATTTTCATACAAGGGCTCTGTGATCTGCTGCCTGGCTCGT
TACCACCCATGATTCTGAATTTCTGCTTCCATGGCATTCTCTAGAATCACCGAGCTTCTCGTTGCAGCTCTGTATCT
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CTTGCCACTTCAGTGCACCAGGCCTGCTCCTACCCTGGGCCATTGTTCTAGGAAAGAGCCTTTCTGGGCTTGACCTCAG
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GGTAGTTCAAGGACATGGCATCAGCATCTGATTAGGATATTTGGGATGCATCATCCCAAACAATGGAAAAATGGAAGGG
CAAGTGAGCACATGCAAAAGAGAAAACATGAGGGGCTGGGCTCACTTTATAACAACCCACTCTTATGATAACTAACCCC
CTCCCACCATAACAGCATTAATTCATTCATGAGTGTGAAGCCCTCCTGACCTAATTACTTCTCACAGGCCCCCACCTCT
TAATACTGTTGCAATGGCAATTAAACATCAACATGGGAAAATACCTTTTTCAATTGACATCCTGCCTCTACCACTTATT
AGTTTTTAGACCTTATGCCCTCTCTGAACCTCAGTTGCCTCTTCTATAAAATGATCATAAAACCAATGTCCCTACCTTC
TTAGAGTTCTAGAGTTCTATGAATAATCTCTACAAAGAGCAGGTGCCCTTGAAAAACATGATTCTGGAACCTAAGGGGG
TGTAATAAACACATGCTTCTTTTACATTGTTTTTTCTTCTCTGCTTTTATTTCCACATTTTTTATTTTCAAGATCT
CCACATGGTTTGATGACGACCACAGCACCTATTCTAGCTATTGGTAACAATTTATTGAGTAACATCATCCATATGCTCC

Fig. 9.298

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CTGTGCCCCAGCATGTTTCATGGGTGTTGTTTTGGGTCACCTCAGGTTGTTTTTCAGATTGTTTCATGAGTGAGCGAAAAGT
TTATTTGCACTGTTCTCAGGGCCATGTCCAGGGAGCTGTCTGTGTTGAATAACAGCATTCACTTTTGATTTAAGAATTT
TATCTAGCTAGATCAAGAAATAGCTGATGGTGACTCTGGAAATCCTTGTCTCAAATTATTCATGCTAAGAAATACCACA
AAAGAACCCTGGGCCCCAGTTTCTCACTTTCTCTAACTGGAATGTTGCACAACTTTCTCTTCCCTGTCCCTCC
AAGACACGTCATGCTGACACCTTGCTCCCATACCCAGGCACGGTGTAGCTGGGGTTGCTGCAGCAACAGCACACAAGG
AGTAACATTTCCCAGTGACTACAGTGGCTGGGATGAGTCAGTTGAACATTTCACTGGGTTTTTATTAGTGTATGATGCA
GTCCCATGTCATCTACAGTTCCTAGCCTGTCCAAGACTTACACGGGATGATGTGGAACACTCCATGCTAATCACCTTCA
AATATTAGATGGTCAAGGGTTCTCTCACTGAAGTCACTGCATTGCTGGTTCCTTCCACAGCACCCCTTACTAATATCTA
CAGTAAGAGGCCAAAAGTAAACTCAGGCCCCACTGCTGGAACACCATTCTACTATTGCATATCCTGTGTGCCTAAAA
AGCAACAGTGACCCTGGCTGATGATCAAGAAGCCCCCTCTTTTTTACCCAGGGAGTTGGACAATTGTTAACATCCCAGAAG
ACATATTTTTTAACCCAGTTGCTGGGAAGAACATGTGCTGAAGTATAAGGATGGAATCACTGTAATTCTCTATGGGAAC
CATGAAAGTGGAACCTTGCCGTTTAATATTGAACGACACTGAGCACAGAGTTATTGATGATTCTAATGCGGGATTGTTGGT
TTATTTTTTCAGACACGGAGATGACTTGATTGTGACTCCATTGCTCAGGTAAGCACAGCTTGGTGAATGGGCAGGTTTC
TCACAGATGTAAAAATTTAATTTGGGGAATTAGTTCGGGTTATTAATTTAATTTAATTTTAAATCAAGCACAAAGTACAA
ATACAAATTTCTTGGTTCATTCAAGCAATTCAAAAGCAATGCTAGAGAAAGTGACTTTGGCTATATTAATCTGTTTCTTA
TAAACCCATAGTACCAAGAGTGCTTTTGCTTCAAATAATAAATTAGCTAAGGGATATACTTAATTGGGCTCATACTGA
CATAAGCACTAGAAGCAGAAGGCTATATCATGGCTTAAGTTATGTTTAGTATAATTAGGTACATGATCAGTGCTTGAAA
TTCAATATTCCAATTCAACCCAATACCTTATTTATCTCTTTTTTACATATGCCATTTGTGCAGAGATTTAGCTGAGTGA
TGTGTTGATGTTTACTTCTTCTTCTAGGCATACAAATCATGAAAATACTTTAACCACATGACATATGTGCAAGCAGGT
GATAACATGATCTAAAATGTTTGGGAATATAAATATGAAAGAGTTGATGACCAGATGTTCTATGTATAGACCAATACTT
CTCAACCCCTACCTGCACGTTAGATTAACCTAGAGACCTCTTATCCCAGTGCCAGGCTGCAGCCCTGGTCAATTAAATC
CAAAAATTTGGAGATCTGTTACAAGCATCTGTATTTGTTAAAGGCCCTTAGTATTTTTCTAGTGTGCAGGAGAAGTTGAG
AGTCTCTGGTATAGACAACTAAGGGCGGAATGAGATAAACTGATTTAACCTGGGGCATGAAGCAGGTTTAAACCAAC
ACTGATGAGTAAGAAATGTTAAATATTAATAATGCAAGTAATCCTGAGAGGTTGTAAAAATTATAATCAGGCCTATATCA
TACATCAAGGTGCTCTTGACCTTGGACAACCTCACTTAAATTTTCTGGGTCCTCTTTTTTTTCACTGGTAGAGTTAGG
GGACTGAGCTGGATGAAAGACAGTCAGATCTAGTTAAACCCTTTCACCGCCTCTTGCCAACTCAATGGACTTGAGCGA
GTCACCTGAACTTTTCCAAGCCTGGTCTCTTCTCTGTTAAATAACAGTTCCAAGAGTACCCACCTCATAAGATTGCTAA
GACACTGAAATAATACCACGCTTATAAAATGCTAGGCACAATTGCTTGAAAATATAATATTATAAAACCTTTGTGATTG
GTTTAAATGTATAGAGCTGAAAATGTTCTTTTCATTTCTAAAATGTCATAAGTATTATTTTTTAAATGGATAATAGGGTT
TAGATAGTAGGATTATGTATAGTTCTACTTTTTTAAACTTTCCAACTTTTATTACTGTAACCTTAGACCTATAATATAA
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CAGGCATGGTGACGGGTGCTGTAAATCCCAGCTACTTGGGAGGCTGAGGCAGGATAATTGCTTGAACCCGGGAGGTGGA
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TCAAGTAATATTTATTAAGCTGCAACAAATAGATTAGTAAATTGGACTTGAACCCTGCCATCAGTGCTCTTAAATTCCA
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TGAGAACTCATTTCTTTATGCCTGAGTCCATCTCCAAAACACGAATCATGGTTTCCAAATCATGGTTTCTTTGACGTG
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CCTGAAACATCTTGGATAGTAATTTTCAGAAAGGATCCTATAGTATCTCTCCATAGTTTATCCATTGTCTAAACACT
GTCAGGGCTAGAAATCTTGGCTGGTGTGTTAACCTAACTCTTTTCCACCTCTGATTTCTTATGCTTTCCCCTGTAAGCT
CAGAATCCCTTTGGTCTCAGAAAAAGTGACAGCTTTAAATATTTTTCTTATTTCAATTGTTAAAGTATTCTCTTGTGT
TTGTGTTGAAATCTTTTGGTGTGGGTGTTTCTTAGCACGAAGGCTAGAGAGGAATCCCACTGGAGTGACGCGGCAAC

Fig. 9.299

TGACCTTTTTTTCTTGGCCAGAGTTCTTGGGGGCCAAGGAGCACTAAGGAGGCACAATGCTGATAAACTGTAGGAACT
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TGAAGCAATTAAAACCTTTGGAGACTCTTGAATTACCAATGCTAGGCATTAAGTAAATTTCCCAAGTGCATGCCAAGTCA
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CAAACCATTTTCAAGTGATTTTAAAATGGCCTTAGCTCAAGAAGTTTCAAGAATCCTTTCTTGTGTAGGTCTTGGCCA
GTCTGCGAACTGTACGAAACAACCTTTGCTGCATTAACCTAATTTGCAAGATCGAGCACCTAGCAAGTAAGATATCCTTTT
TTCCCCCTCTTTTTTGTGCTGAACCTTTTAAATAAGTTTTCTCACTTGATTGAGCCCAAGGCAATATGTTAGACTTCTCT
GTGTTGTTGCCATGTTGTCTGTGTATACTAAGTCATATGATGTCCTGTTAATTTTCTATAAATACTTCGTGGTGATGGT
TCTAATATCAGAAATGAAGCAGTATGACAAATAAATATGGTGATTCCATCTGTCAGAAATCACCTGGCATGATCAGTCC
TCCGCCAGTTATTTACACTCAGGGTAACTTTATAGTTTCCGGCTTACTTCATAAATTAACCGTGGGTGAATAATCTCAT

Fig. 9.300

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CACACCTTAAGATCAGCTGAAAGTCATCCTCTCAAGACAACGATGATTTGGAGAAGCGCCATATCATAGCAAAGATCTT
TGATTCTCCTGCTGTTACTGACTTTCAGATCAGCAGCATTTCATGGAGCAATTAATAGAATTGTGGTTTATATGACAACA
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CCTTGTAATTGATTCTATGTTAGCACGTAGCTCTCTGTGCTCCTGTTGCCCTGTTTGTCTGTTGGGATGCTCCTGAACA
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AAAATAATTTCTTTAAAGTAAAATGCTATATCTGAAATGACTCAGAGAGTGATCAAATCAATAAGCCCCACATATTTACT
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AGAAAGATCTGGAGCTTATTCTTCATGTGTCTAGGAAGAAACCATTTCTGCCAAGAGTCAATATAACACCAACACCAAT
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TTATACTCATTTGAGCATTTGCTCTGGGCATTTGTATTTTGAGAGATGATGTTACTTTCAAGTCACTTCCCATTTCTGGTT
CCACACATGTGTTATTAGCATGATATAATATGTCATCATTTAGTTTAATAAGAGTGAAGGCATAAATAAAAGGAGTAAC
GTGTTGGTTTCATCAATATCCCCTTGTGTGTAGGAAGCAACATTGTACCAGTAGTCCAATAAAGGACTGAGAGAGCTGA
GAATAAACTCAGTTTGGGAGGGGAGAGAAGTTAATATACAATGCAGAAAAGTTCAATGAGATCAAAACGTTGGGATCT

Fig. 9.301

ATAGATGTTTGGGAAAGAAAAATAGGAGGAAAGATGAGAGGAGCCAAGTGCATAAAAAATAAAATTATCTCAATTCCTAT
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GGTGATGACTTCCAATTATCATTTTATAGTACATATATGGTTAACCAGTTTTGTTCTTGATATTGATCAAGAGATGAAC
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GGGACACAGAAATACCTTATCTAGAAAGGGAACTAATTAAGTGTGTTACCACTTAGTGCTGCTGGGCTCCTCACAAA
GCACAACAGGAAATGAACTGACTAAACTTGGGAAGTCGTTTTTCTTGCCCAAGCATAGCAAACCAAGGTGGGAACATGG
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CCACCTTGGCCTCCCAAATGCTGGGATTACAGGCATGAGCCACCACACCTGGCCGAGAGTTTGATTTTATAGCATTAG
GGTTTTAAAGCTAGGTTTTAAAGGTAGGTCACTGGCCAGTTTTTATTTCAATATATAGTAGGTAAACATACAGGTCTAA
AATGATCTAACAATTCCTTAAAGTAAGGCTTTGAAGTTTGCAATTTATAAAAGAGACTTAAATAGCTCTTTTGCTCTTT
AGTGTGATATGACAAAGATGATGTGTGGCATTTGGAGCCTGAATGTGAACCCAGTCTCTCCTTTTTGCTTCATTTCTC
ATTTTGGAGATTTGAGTACCTAACACTTAGGACTGTTGCAAGAATTCAAGGAGATAAGTTATATAAAAGGATAGAGTTC
AAGTTGGGCATGGTGGCACACCCCTATAATCCCGGATATTCAAGGAGACCAAGGTGGGAGGACTGCTTGAGATCAGGAAT
TCAAGACCAGCCTGGGCAATGTAATGAGACCTGTCTCATAAAGTAAAAAATAAAAAAAAAAAAAAAAAAATGGAGCCGGG
CATGGTGGCTCATGCCTGTAATCCAGCACTTTGGGAGCCCAAGGCGGGTGGATCACCAGGTGAGGATTCAGGAGTTCAAGACCAG
CCTGGCCAAGATGGTGAAACCCCGTCTCTATTAATAACAAAATTAGCCAGGCGTGGTGGCAGGTGCCTGTAATCCCA
GCTACTCGGGAGGCTGAGACAGGAAATCGCTTGAACCTTGAGGGCGGAGGTTGCAGTGAGCCGAGATCAGGCCACTGCT
CTTCGGCCTGGGTGACAGAGTGAGACACCGTCTCAAAAAAAAAAAAAAAAAAAAAAAAAAAGAAAGAAAATGGATAG
AGTTCCGAATGTGATGAGGAACTACCAAGTGTTCCTTCTATAACCCTGTCCCAATGCCTGACATTTTTTTTGTTCCTA
AAGCTAATTTAAAACTGGTAGAATAGAGGTAACAGAGAATATTGGTATGTCAGCTTCTTTGGTACATTTTATTTATGA
TTTTGTTTTTACACAGTCATGGCATGACATTTTTTAGTAATCCTTTTATCATTTAGAGTAAGGCTCACCTACATACGTCC

Fig. 9.302

TTAATGTGTGGTGCACACCAGAAGTAAAGAGTGCAATTGGAACATGAGAGGGTGGCAAAGAGTCAGAGATGCCCCAAGCCA
TACCTGCACTGGGACAGTGATCCCATTTTTGAAGTACAGCCATGTAGCTTGGAGCACAGTCCTTCTTGTAAGCATTTC
ATTTTCGAACAACCTTAGCTCAAGGCAAGCATTCTGCAGCCATCTCATTAATGTTACATCTTCACAATACCATTTTCA
ATGCTTCTGAATTTCCATATTTCCAATATGTTTATTTAATCATCCTCAATTAGAGGATTTTCATGCAAGGACACAACCTC
TAAAGAAAACGTATTTAAATATCTTTATTTTAAAGATGAAAAAAGTACCTAGTGAGAGAGGGTGCTGCTTAGTGAC
AATGCCAGGAATAAACATGTGGTTCCTTTTTTCGAGTCCATGGTTCCTTCTTACACCTTTATAAAGTAAATAAACCATTA
GATTCGGGTGGCCTTTATACCCACCGTGGAAAGTATATGCTTTAGAGAAAGTAAATGAAGTATTAATACTAACTCCTGAG
TAATAGATGTGGCATTGTATTATTTGTATATGAGCTTGAAAATTTCTCATCCCAACCTCTCACTGGACATATCTCTGTA
GTTATAGCTATTTTGATTATATCTACTGCTTATACCCAGAGGCATAGAGACCAGCACCCCTAAAGCAAAAGACTTCTGG
ATTTTGACTTTGGAAAATACAAATTCCTTCTACAGGAGCGGTGAAACTAGTACAGCTGCCTGCTCTTTACCTGTAGGGGG
AGACACATGATGCAGTGACAAGCATAGAACAGCACAGAAGTGGTTTTAGTCCAGAAAATAAATAAATAAGTAAATTGGT
CTAAGTCTTCATTCTTTTTGTATTTTTTTACCCTGAAGATTCATGGCTTCATTAATAAACAGTAAGTGCATTCTAAATA
GAATTTAGTCAATTGGCACGTAGAAAATTAATAAGTAAACTATCCAAATAATGCTAGTCTGTCAAACACAATGAGAAG
AAATTTAATCTTATATGTTTGTGTCTTAAAGCACAGCAGTTTCTAAGAATTAATAAGGGGGAAAAACAATAATGATA
TTAATTGGGACAATTATTTTTTTTGTATCTGGGCTAATCACTTAATGCTTTGAGCTTATCGAATGAAAGAACTAAGC
AACTATTTTCATGGTGACAATAAGGATGCTGGCTACTTACTTAGATCTGCCTTTTAAATCTGTGTGTGTGTGATAAAAGC
AAGTAGTATATATAGTTTTTTTAAAGAAATATAGAGAGATCTAATAGATGGGAGCAAGAACCATATAAAGAAGGCCAAA
AGGGATCTACATGAGAGAAATATGTAAGTCCTCAGGAGATACTGAAGTAAACATGTGTTGTGTGTCAGTGTATAGGGAAG
ACAAGAGACCCAGCCATTGCCAGAGAAGCCCTACTTCCCTCATCAACCACCAGCAGGCAGCAGCTTCATGCTGTAGTTCT
TTTTTTCCTTTAATAGAATAAAGATTGTGGCCACACTTTAATAATTGCCAGAAATATGTGTTTCATTAAGCTTTTTTAAGA
AATGTGATCGAAAAATGCTGATGATACATTGTGTTTAAATAGCTTTAATAAAAAAAGTATAAAGCATCTAATACATCTAA
TGACTGTGCGATTTCCTAAATCTTTTAAAGCTGAAGGAGTTTACTTTTACAAAAATCCTCATTGAAAAAGAATCTTCCGG
TAAAAACATTGAACTGAAAGATTTGCATAAATGCATGGAGTATTACTATTTATTAGTTAAACTTATGATCTTTTAGAGC
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ATTAATATATAAAAAGATCAGATCTGACACCGTGGCTTTTTGACCAGGAATTGATTGAAGTTAGACTTGTTTTAAGAAT
AGTGAAAACAAGGGCTATGTCCTAATTATGAGGGGAAATAAATTAATTGTGTCTCTCTTGTCTGAATATCTGATTTGGA
TGCTTTGTTATAACTTTAATTGAAATAATTAATGGGGCTCATGTTTCATTTAGCCTAGCAAAAAGGTGATGAGCTAGCA
TATATGTTAAAAACCCAGCTTTTCTCTTAGTAGTTGTCTAATTTTTGGTCAATGTACTTAAATTCTCCAAGTTTTGGGGG
TATTCTTTATCAGTAAATTCTAATTCAGTGCTTAACATAGTACTTGGTACTCAGAATGTATTCAACAAATGTTACTTAT
CATTATTGCTATAATATTATTATTATTATCTGCTTGGTCAACAGTATTTTCCATGCACCTTATATATGGCATGCAT
ACGCTCTGGAGGTAAATGTGACTACTGGTTCAAGAAATTTACAAAGTAAGGGAGAAGGAATTATAAATAGTTAAGTAC
TAGCTAAATGCCATGATTGAAGCCTTTATATTTTTTTTATTTTAAATTAATTAATTTTATAGAAATAGGGCCTCACTCT
GTTGCCCAGGCTGGAGTGAAGTGGTACGATCTTGTCTCATTTGCAACCTCCGCCTCCAGTCTCCAGCAATCCTCCCACC
TCAGCCTCCTGAGTAGCTGGGACTACAGGCATGCACCACCACGTGTGCATAATTTTTTATTTTTTATTTTTTGTATTTTTT
GTAGAGGCGGCGTTTTGCGGTGTTGCCCAGGCTGATCTGGAACCTCCTAGGCTCAAGCAATCTGCCCACCTTGGCCTCCC
AAAGTGCTGGGATTATAGGCATGAGCCACAGAGCCTGGACTGAAACCTCTTTAAATATATATATTTATATAAATATATA
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ATATGCTTATTATATAAATAAATATATTTCTCATATAAATATATATGTTTATTATAAATAAATAAATAAATAAATATATT
AAATAATTATATTATATATTTAATATATAATATATATGAATATGTATTCATATATAGAGAGAGAGAGAGCCAGTGGCTC
TACCTAGGAAATGCTCCCTGATAAATTTGAAGAACATAATGCTTTCCATGTGGTTAGGGGAGGATTGAGGATTACCTCA
CAACAAAAGACCAGCAAGAATAAGGTATGGAAGCATGGAATGCCTGTCCATATCAGGGGACTGGGAACAGTAAAGGACA
AATTGAGTTAAAGGTGTATAGGAGCAACTGAACAGGATGCTCCAGAGGTAGATTAAGGAAATAATTTGAGGCCTACA
TAAATAACAGTGGACTTTATTTTTATGGACAGTGGAAAATCATTGAGCATTTCGAATCGTAGGAACAAATACACATTTCA
GAATGACAGTAATTGGAAGCAGTGTAGAAAGAGATTGAAAAGAAGAAAGACTAGGGGCAGCCCCATACTCTTCAGGGGG
CTGTCACAGGCCTTGTACAGGATCAAGATAATGCAGCTCTGAACTGGGGTGGAAAGTGGGGCTGTGGAGATGAGGGAATG
GGTTCACAACTCTTCAGAGATAGGATTGAAGGGACTTGGGGGTTGATTAGAGCAGAGACTGAGGATGAACAATTTTAAA
GTGAAAGTATGAGCCAGGTGTGGTGGCTCACGCCGTGAATCCCAGCAGTTTGGGAGGCTGAGGCGGGTGGATCACCTGA
GGTCAGGAGTTCAAGACCAGCCTGGCCAACATGGTGAACCCCATCTCTACTAAAAATACAAAAATTAGCCAGGCGTGG
TGGCACGTGCTTGTAAATCCCAGCTACTCTGGAGGCTGAGGCAGGAGAATTGCTTGAACCCAGGAGGCAGAGGTTACAGT
GAGCCGAGATGGCACCATTGCACTCTAGCCTGGGCAACAAGAGAGAACTTCATCTCAAAAAAAAAAAAAAAAAAAGTGAA
AGTATATATACACATGTTTCATTACAGCACTGTGAATGTGCATAAAGGGTTAAAAACAATCCAAATTCATAAATAATGTGT
GAATTCATTTAGGTAGGGGTAAAAAATTTCCATAATAAATGATTTTAAATAATTTATTTTTGAGAAAATTTTAATA
ATATGGAAAATGTTTGTATTTTAAAGGAATACATAATAAAGTGCCATGTGGGCAATAGGCACACATTTAATGCATTTA
ATTATGTATATTGAATTGAACTTAAGTTACTTTTAGTTGTTGACCAGTAGTTAGCAAAGGTAGATAGGTTTACCATTTT
ACATTGAATCTAGTGACAAACATGTTATTTCTCAGGTCCCAGTTGTTAGTTTGCCTCTCCTTGCCTAGAAAGGGCACTG
GTGAGACCCGCCCTAACAATATTTCAATATTATGGTTCCACAGTCCAGCATTAAGTGTATTTTAAATAAATAATATCTC
TCACATAGGGCAGCATTTTATGGATTATTGAGCCTATGACAAAAATGTTGTATTATCCTTCAGATGATCAGAAAACTA
CCCTTAGGGCTGGGTGCAGTGGCTCACACCTGTAATTCCAGCACTTTGGGAGGCAAGGGAGGAAGATGGCTTGAGGCCA

Fig. 9.303

Fig. 9.304

TGTTGCACACCTGTAATCCCAGCTCCTTGGGAGGCTGAGGTAGGAGAATCACTTGAACCTGGGAGGCAGAGGTTGCAGT
AAGCTGAGATCGTGCCACTGCACTCCAGCCTGGATAACAGAGGAAGACTCTGTCTTAGGAAAAAAGATATTTATGA
AAAATGAAGCAAAATGCATGGCTCACAGTAAGCACTTGATCATTGCTGGTTAACTATTATTAAGATTATCTAATAGGGA
TACAAAGTATATTCCTGAAAGAAATTAGGGTGCTCTTAGATAGGGGAAATAAATGCTAGTAGGATTGGTTTCAATGCAC
TATTTATTTTATTAATTTATTTCTCACTGAATTTCTTTTTTATGATACTCATTTTTTTTAGAGATCCAGATGGTACCTGT
CATTAATGTATCCCTGAGAACTCAGTTCCTCTGAAGGCACAAAAATAATTTGACCATCTCATCATTGATCCACACACA
TTTATTGATGCCTCTTTTATGCTTGGTTCTATACAAGACGCCGTGACGTGTGGCCCCCTGTAGGAAGTCCTGGCCTCTGT
CTCTAAATCTGTCATCTCAGCCCAACCCCTATTAGATTATCTGTGCCCTCTGCGTAGCCATTCACTTCGCCTGGGTGC
AATCATGCCTTCCTTACCCAACGTTTAAACAAGTCCAGCCTAGCAACCTACATCATGCCATCCAACAGAAGGAAGGCATT
ATTGACTCAGAACAAAGTATAGGCGCTCTTAGAATAAGTTTATTAAATCATTTTTTTTAAAAAAGATATGTGCTTTTT
CCCAGGGGAGCAGTGAGGCATTGTGAGGGGTGGATTCAATAATCAATACAGAGAATCCTGGCTGCACTATTTGTTAG
CTATTTGACTTTGGGAAAGTTATTTCTCTGAGCCTTGATTTCTCATTAGAAGACGGGAATAATAGCAAACCTATCTCAGA
GCATTAAGTAAGAATTAAATGAAGTGAATTAAGCACGAAACCTAGCATCTGACGCATAGTAGGAATAAAAAAGAAATA
CTAGTTCTTGAATAAACATTTTTTTCTTGAAGTGTACTGTGATACTTCCATCAGTGAGTTTCCAAGGCAAATGAATCATC
TTTAGAATTGGAAGCTCAATGTAAGCAACAATGAAGTAAAGAAAACAACTGAATTTCTCTTGAAATTATTTTCTTTAC
TGACTTGTCTTTATTGTCTGTGTTTACAAGAACTAGCTCTGACTGCAAAAATGTCTGCCTGTTTCTATGGCCAGTTCT
TTTCTACATGTTTAAATTCCTAATTGTGTCACACAATCTTAGTCACAATATTTTCATGTGCAAGCAATTAAGACTCATTT
AACAGCCCTCCCTCTGGAAGGTTTTGTTCTCCATCTGCCACAAATCAGATTCCCTTAGGAAGATATTTGATTTTGAAACA
ATGTTAAAGTACTGTTTCAATTTTCAATTTTCAATTTATGTACCATTTTTTAAAGTGATGTAAATGGACAGCCACAAAA
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ATTTAAATATGGGAAATAGGAATATACCCAGTTGCCACTCTGAATCTTAGCTGTCCTGAGTTCACTGCAATGTATA
TAATAAGGAAGTCAGAAGGTGGAAAAGAGGAAAAGATTAAATTTGTAAGTTTTTTTGAAAGATATATGGTAAAAAGTAGA
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GGAGGCCAAGGAGGGGCGGATCACCTGAGGTGAGGAGTTCGAGACCAGCCTGGCCAACATGGAGAAACCCCGTCTCTAT
TAAAAATACAAAATTAGCCAGGCGTGGTGGCACATGCCTGTAATCCCGGCTACTCGGGAGACTGAGGCAGGAGAATCGC
TTGAACCTGGGAGGCAGAGGTGTGGTGAGCTGAGATCGCGCCATTGCACTTCAGCCTAGGCAAAAAGTGAACTGTCT
CAAAAAAAGATTTGTTTCAATTTAAATCAATTCGATTGGCCAGGCACGGTGGCTCACACCTGTAATCCCA
CAGCACTTTGGGTGGCCGAGGCAGGCGGATCGATTGAGGCCAGGAGTTCAAGATCAGCCTGGGCAACATGGTGAAACCC
CCGTCTCTACTGAAAAAAGTAGCCAGGCATGGAGGCATGGTGGCACATGCCTGTAATCCAGCTACTTG
GGAAGCTGAGGCACGAAAATGGCTTGAACCCGGGAGGCAGAGGTGAGATCATGCCACTGTACTCCAACCTGGGTGACAG
AGCAAGACTCTGCCTCGAAGAAAAAATCAATTAGATAAGTGAGAGTGTATATTCAGGGCACTTAAATCTATG
CTCTCAGATTAAAAATAAGATTTAATTACAATTTTTTTTTTTTGTATGCAGGGTCTCACTTTTGTCTCAGGCTAGATT
GCGGTGGCACAAATCACAGCTCACTGCAGCCTCGACCTCCTGGGCTCAAGCAATCCTCCACCTCATCCTCTGTCTACC
TGGGACTACAGGCACATAGCACTGCACCACCATACTGGCTAATTTTTTAATTTTTTTTTTTTTTTTAGAGACAGGATC
TCACGATGTTGCCAGGCTAGTCTTGCACTCCTGGGCTCAAGTGATTCTCCTGCCTTGGCCTCCCAAGTGCTGGGATT
ACAGGTGTGGGCCATCACACCCAGCCAATTTTGAAGTATTTAATTTTTTAAATGAAAAATTGATTAGATAATAGTTCTC
CCTCACTACAGGTGAAGTCTGTTTTTATTTATTTGTTCAATGGGCTTCTTTAGAACATGACATAGAAGGCAATCCTTGG
TCAATTAAGGCAGAAACAAGAATTTATTAGGTTCTGAACATAAATACTGTCTGTGAACTGGTAACTCTCTAATTAAG
CATAAATGTGAAAAGAAGAGGATTAGCTCTTCTTGAGGAGTTGGAAATGGAAAATATTACAATTTGGAGAGGTAGCTTG
CAGAAACCGTACAGTTTTCTCTGCTTATATGCCCCAGCGTTGGGAGACTTGAAAGGAATCACCACCAAGTTAATGCAA
TAAATTTCCATATATAGATCAATTTGGATGTTTTGTCCCCAGCTTCCTAGGCCTTTAATAAACTGAATTGTTTTGGTATC
ACTGGATGAAAGGTTCTGTAAAAGTTCAAAGTATTGTTATTTGGGGCATTACACCTGCATGTTTAAATGCCTTTGTG
CAGAAATGTAGTCCAATCTGCTTTCAAATCAGACCTTCAAAGAAGGGTAGATTTCATCCATGTGACAGATCCCCCTTAGGTG
CTTCTCTAAAAGAAAAGTTAAGCTAGCAGGTCTACCCCACTCATTATCTTTGTCTCTTTGTGCGTAATCATCAAACC
GCACCTCTGAAAAGAAGACCAAGAGAAAACCTTTAGTATCTCTTTGCTGGAGATGCAAAGCAAGATATAGAAGGAAGTGG
AAAATAGTTCTTAAGTAAATACAGGGGGGAAAACGGAAATTGAGAGGACGTACTTTTCGCTAACAGTTGTAACTAAAA
TAAATTTGAAGCACGCCTGCACCCTCACCTGAATGGACTTCCTCCTTGGCCAGGGCACTTTAAATTTAACCTGAAAG
ACTGATTTAGGCCGCAAAGGAAGTCAGACATGCCTTATTTTACCCCTCCAGTATTAACATCACACAGACCTTAAGTCT
GATAAGAAACATTTAGGATCTCTTTTCTTGGAAGCCTGCTACCTGGAGGCTTCATCTGCCTAATAAACCTTTGGTCTCC
ACAACCTTTTATCTTAACCCAGACATTCCTTTCTACTGATAATACTCTTTCAACCAATTGCTAATCAGAATATGTTGAA
ATCTACCTGTGACCTCGAAGCCCTCCCCCACTTTGAGTTTTCCCGCTTTCCAGCTTTCCAGATAGAACCAGTGTAAT
CTTACATGTATTGATTGATGTATTATTTCTTCTTAAATGTACAAAAACAAGCTGTAGCCTGACCACCTTGGGCACATG
TCTTCAGGACCACCTGAGGCAGTGTACACGTGCATCCTTAACCTTTGGCAAAATACACTTTCTAACTGATTGAGACCT
GTCTCAGATATTTTGGGCTAACACAATGAATATGAAAAAATTTTTTGTGCGGGGGTGGGAGGTGGGACGGAGTCTC
TCTCTGTTGCCAGGCTGGAGTGCAATGGCGCAATTCGGCTCACTGCAACCTCCACCTCCTGGATTCAAGTGATTCTCC
TGCTCAGCCTCCTGAGTAGCTGGGATTACAGGCACACGTACCATGCCTGGCTAATTTTTTGCAATTTTGTAGTAGAGACA
GGGTTTCACCATGTTGGTCAGGCTGGTCTTGAACCTCTGACCTCATGATCCACCCATCTCGGCCTCCCAAAGTGCTGGG
ATTACAGGGCGTGAGCCACCACACCCGGCCAAAAACAATTTTTTAAAGAGCATCTAAGCTCAGAAATCACAGGCATA
TTACGCCTGCCACTAAGGGAGTTAGTTCCATTGAAGGATATAATTAAGAGTGAAATGAATGGTGTGCTAAGCACTTAGG
GATAGTGGTTCACAAATTTGTCTGTTCTTAAATCACCTTGGGGGGCAGGCGTGGTGCCTCACACCTGTAATCCCAGC

Fig. 9.305

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ACTTTAGCAGGTCAAGGCCAGAGGATCACTTGAGGCCAAGAGTTCAGACTAGCCTGGGCAGCGCAGTGAGATCCTGTT
TCTACAGAAGATAATAATAATCACCTGGAAATTTCTTTTAAATGCAGCTAATGAAGGAAAAAAATCCGAGGTTT
TGAAACCAACTCAAAGACAACAGCGATATTCAAGCATAAGATGTAATAAAGGTTGTACACTAGATAGCTAGATAGCCAG
AATAAAAAGGAAGAGATAGTTACAAAAAATATAAGGAGGATAAATGTATAGGATTTTCATAACTGCTAATCATATGATTT
TACTGAGTAGGTGTAAAATGCTTCTGATAATGTGTGAAAATTTATAATCCTTCGTATTATATGTAGGATAAACATAGGT
TAAGACCTGGATTCTAAGGCTGAATTTAAGGCTAGTTTATCTCCATCCTTAGATTTCTTACATTTTCATTTAAGAGAAAA
TGTCCTGTATATTGAATATTCATGAAAATCTCTGAAAGGTGTTATGCTTATTCTTAACCTCTTAAAGGTGTACACTGAA
TGTAATTAAATCATTTTTGCTGGCTCTGGTTCCTCATGAACATCTGCTTTTGTACTTCCCTGTCATTACAAATGCACT
TAGGAGCTAATGATCTATGAGGACTTTTTTTTCCCCTACAGTAACGAGCAGCAAATCTGGCTGCACTTTAATTTCTCAT
CTGCTGTCCCATATTGTCTGGTGGTCAGTTCATGATGTTACTAAGCTTGGCTTTATTGGCATCTTTTGTGAGCTGCTGC
TGCTTTTTCTTGGCAAACCTGATTACCAAGCTATTGCATTGAGCATAAAATAAAGGTAAATTAATTCAAATAAAAGT
GAAGGTTGAGGGCAATTCATTTCTGAGGTAGACCTTTAGGATATGAGATGCATAAAGTGAACAAGATCCTACAAGTGTT
TACTTGACTTTTCTGGGTCTTTTTCACCTACTTACTGATTTTGAATAGTATAAATTCCTGGATAATTAATCTGGATAAG
TAAGTCGTCACTGTACCTCTAGAGAAAATAAAATCAACCAAATATGTTTAAATCTGTGCTCTGGGTTCAGAAAACA
AAAATGAATAAGATATAGTCTACCCCCAAGGACTTGACACAATATAATTGTACATGTGCAAAAGAACTGTCTAGGTG
TGGTGGCTCACACCTCTAATCGCAGCACTTTGGGAGGCTGAGGCAGGAGGATCACTTGAGCCTAGGAATTCAGACCAG
CCTGGGCAACATGATGAAACCTTTTCTCTACAAAAAAAATAACAAAATTAGCCAGGCATGGTGGCACATGCCTGTAG
TCCTAGCTACTGGGGGAGTTGAGATAGGAGGGTGTCTTGAGCCCAAGAGGTCTAGGCTGCAATGAGCTATGATCACAGC
ACTCCAGCCTGGGCAAAAATCTTAAGTAGTCTCAGGACTGTACCACAGAGTATCGTAAGAATTCAGAGGAGGCAAGA
CCAAATTAGAATAATAAGCACATGAAGGCTTCCAGCAGAACATGGTATTTTTGTTGGGCCTTGAACACTCTTTAGATGC
TTTAGTTTAAATGTGCCATAGTCACACTTTCTGTATTGGGAGTGTAAATGGGTGATAACTACTCCAGAGCTTTAGGATTG
CTTCCAGTATCCCAGCAAAGCAGCCCTTTTCAACTAGAACCGTTTGCTATTACAAAAGAGAGGTGATCACTTGTGATTT
CTTAACATTTCTTCACTTTGCCTCTGGCACTGGGCTTCTGAAAGTCCAGGAAAGAGCAATGACCTCAGGGTTTTAAGAC
CAGGGGTATAATCCCAGCTCTGCCTAGCTCTCTGTGTAATTTTTGGTTCAGTAATTTAACCTGGGTGTGTTCTGTAAAA
TGATGATATTGGATTAAATAGTACCTAATCATCTAATTTTTCTTAATATTTTATTATTAATAAAAAAATGCATGCCTCT
GATGAATTGCCTTTATCTTCTTTAGTCATATTTCCCTAAGAAGTGAATAAAAGATAACCAAGGCAATGTGTGAATTCAC
TTTTTCCAATCTGGATGTTTAGGGGATATCCTTGACACCATTTGCTATTTTGAGTTTTCAACAAAGAGTTAAAAGAAAA
TTCTGGCACTCCTATCTAGTCATCCTCTCCAGTTGGCAGAAGTCTTCATGTGGACTTGATGGTTGCCCAGAGCAACAAA
ATATTAGGGACAGAAACATGTTCAAGGACTCGATTGTATAAGTGAAGTCAAGAGCTGAGAGACCTTTTCCAGCTTGACTGC
AGCCCACTAGCTAAAGTGGGTATTTGTCTATTCTGTCTGCATACTGTGACTTGAGAGATGCCTATTATTTTGCTTG
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CCCAATGGAGAGCCACTCACGCAAATCAATACCCTTTCTCTCAGTTGGAGCCAGACATCTCTAATCTCTCTGA
ATTAAACTGGATTTTTAAATGTTTTTCTGTCCCTAACTCTTTAAATCCTGAGGTCATCCTTTCTTTGTTCCATCAGT
CTGGCTCCATGATACCAATTACTCCAGCTTGTTAAAGCAGTTATTGGCATATGGTAGTCATGTCTTTTGTCTTATGCA
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CCTCTTGGTTTGGACATTATWATTTGAAATGAATATTCTTTTTAAATGATTGGAACTTAGTCGTAAATTCAAGTGGTT
TACAATAGTAATCTTATCCCAGFAACCACAGCACCTGTTTAGAAAAATGTCTTCGGATCACTTGTTTGCAAATGTCTT
TTTCTTAGGATCCTGGATGGAATTGAACCCATATACGTTACTTGACATGTGAAACACGTGTGACCCTGGCAGATGATT
TGGCTGACCTTGAAACTACAGCTGTTTAGTCACTTTGAAAACAATGCAATACAAGTGATTTACTAGGCTTCAGTTTAA
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TAGGCATTACATGGTAGAAAGATAGCATTTCTGTTCCAAGAAATCTCTATTCTGTTCTAGTTGCTGTGTATATATTCTC
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TTTACTAGATTCATAGGGCTTTTGTCTTCATGGTCTGAGATGACTACAAGAGCCCTAGCCATCACAACTAAGTGGCCAG
CAGCAAGAAGGAGGAAAAAGAAGGAATAGTACCCCACTCCTTTTAGAAAAGGCTTCTGAGAAATCCCACATAATGCT
TCTACTTACATCTCCTTGGCTCAAATTTAGTGTCTGCCACAAAAGAGAATAAGACACACACACACACACACACACAC
ACACACACACACACACATATATATATATTTTTTTTCTGGCAAGTCTCTAGTTTCAAGGATTTCAATCTCCTTTA
AGCAAATGAGTCTTCTCGCTAACCTCTCCCTCTGGCCTGCCAGTCTCTCTCGAATCTCATTGCCATCCTGTCTCTCTT
ACTCACTTTGGGAAAATCCAGCTCTGTGAGTTTTACAGTCTGCCTTTTCCACTTCTTACCTGAATTCCTGGGAAAAAT
AATCCCACATCACTTCAAGCTGGAATTTCAACAAAGTCAAGACCTGTACCTCATCGGACTCTCAGAGCACTTGACAGT
TAGTTCCTGCTGTGCCCTCTCTCAGTCTCTTTCATGCTAATTCCAAACCTGTCTCTCTCTTAAGCCTCCCAAACAG
CAGCTCTTCTCTCTCTTTCAGCAAATAACTTGTGTGCTTTTTGAAGAAAGTCTGTGTCAAGGCATGAATTCCTCAGCTC
CTACATCCTCTTGCCATAGATGTTATCCATAGGTCTCTTACTGACATCCATAATCCCTTCTCTCTCTGCTCTCAGCG

Fig. 9.306

GATGACGTTTCCCTTCTCCCTCCTGCTCTAAGTCAACCCCTGCATCTGGGCCCTTGATCCTCTCTCCTCCCAGCTCCCA
CAAGTCCTTGCTTGGTCTGTTTTCTTTCTATCTCTCACATCTTCAATCTTTACTTTTATTCTGGCTCTTTCCCCATTC
ATGTCTCTCACAATCTTAAAAACAATACACAGCAAACCACAGCAAAGTTCAGCCAACTCTTTATTGATCCTTTAGCTAG
TTCTTGATCTTCTCTTTTCAGTCAAGTTGCTCTACCTCCTCATCCCTATTGAGTCTGCAGTCCACTGTAGTACGACTTT
CTCACAACACCTCTGTTCCAGCTGCCCTCCCTTGGTTGCCAATGAACTCCTAATTGTCTTCTTAGTTCTTAGAGGCTTC
TTCTTCTTAAACTCTTTCTTTGTTTCATTTACTGAGTAAATATTTATTCAACATGTTTTTGTGTTGGGAGCCATTTTGTTC
AGATGTTTCCACACTCTTCTTGCTGCTCTGAATACTTCTTTTCACTATTCTTCATGGTCTCTCTTCCCTCCCCACTTTAC
CTCTTAAATGTGAGTATTCTAGAATGTTCTTTTGTATTAAATTGGTTTTCTCACTCTGTGTACTCTGTAGATGTTTTCAT
GTACCTCATAGTTGTATTTTCCATCTTATGAAAGTGGGTCCCAAACCTCTGTCTCTACTCAGAGTAGGCACCTTCCCCAA
TTTCCAAATTTATTACACAAAAGCCCAATGTCCATTGATATCCAACCTCTTCAAATCCAATATGTCCAGAGCTGGACTC
ACATATTTCTTTGCAAATCTGTTCTTCTCTATCTCAGTTAGTGAAACACTTTTAGCCAAGATCGAAACTTATAAATCATG
AATCCTTCTACCAAATCTATCTGTCTGTCTATATATATTTTGAACAGAGTTGTCTCACTCTGTTGCCCAGGATGGAG
TGCAGTGGCAAGATTTAAGCTCACTGTAACCTCCACCTCCTGGGTTCAAGCAATTCTTGTGCCTCAGCCTCCCAAGTAG
CTGGTATTACAGGTGTACACCATCACACCCAGCTAATTTTTGTATTTTGTAGTAGAGACAGCGTTTCACTATGTTGGCCA
GGCTGGTCTCGAATTCTTGGCCTCAAGTGATCCTCCTGCCTTGGCCTCCAAAGTGCTGAGGTTACAGGTGTGAGCCACC
ATGCTCGGCCCCCAATGAATACAATTTTTATTTTATTTTATAACTATTGATCAAATCCATGCTCTCTTCTCTAACCTGC
TAATATTTCCCTAGTTCAAGCACTACCAACTCTCCCTGAATTATAGTACTAGTACCTGAAATAGTGTTTTTGAACCT
GTCAGTTGCATTTAGCATTTTATTTTAAAAATATGATACAGAATTGCGTAAAAAATAAGAGTATAAAACCACATCTATA
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AAGCCTTCCCATCATCCATGATAAAATCAGATCTCTTTCGTGTGGTACCCAAAGCCCCCTATCCAGATATGTATTGTACC
TCCTCTCCAGCCCAGCCCCCTGGCTGACTCTTTCAGCAGCAAACCTTTTTCCACCCACGTGCACCATGTTTCACAACTGTC
TTCATTTACTTGTACTCTTCTCACCATCTAAACTCGCTTCTCTCTCATAATTGCCTGGTTCACFTTATCTCATTTTAG
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TAATGAAATTGCTCTAAAAATTATGGTCTATGCCTCTTTATTAAAAAACAACAACAACAAAAAACCCTGCC
CACCAGTATAGCAATTCTTTTGTATATAACCACACATAGTACTAGTGTGTGGTTCTGGGGCACTGTCATGCCAACTTACC
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CACAGAGTAAGCATCTTTTCTTTACATAGTAATTCACAAACGTCCCTCATCACCATATGACAATATCCTCACTGGATCA
GCTCGGTTACCAGAAATAACTAGATCAAATAAATGTCATTCTCACATGGACACAGGTGGACTAGGGTTTAGAAGTTTT
GGGAACCATCACAGTTCTAGGCACCTGAAACTTCTTCTTTTGTGTTGTAGTAAATATGGTGTCTTTCTCAGGACTCTGCC

Fig. 9.307

CTTTGTTTATTTATATATCTGGTCTGGTTTTTGTCTTCTAGAGATAGTTCTCTACTCTCCATAGTGTCTGCTTTCTCAT
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TGTCCATAGTCTTAACTTTGTTAATAACAGACTGCATGCCCTGGAAAACTATTTCACTTCTCTTGGCCTCATCAACA

Fig. 9.309

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GTGAAACCTGTCTCTACTAAAAATACAAAAAATTAGCCACGCGTGGTGGCGGGCGCCTGTAGTCCCAGCTACTAGGGA
GGCTGAGGCAGGAGAATGGCGTGAACCCGGGAGGCGGAGCTTGCAGCGAGCTGAGATCTCGCCACTGCACTCCAGCCTG
GGCGACAGAGCGAGACTCCATCTCAAGAAAAAAGAAAAAAGAACTTTCCCTCTTAAAAATATTCTTGTTTAAATCT
AAATTAATTTTCATAGATACCATTTCCATTTTTCTAATGAATAATACTGTATCCTTTCCCTATCTATACCATGAAGTTTTT
CTTTGCCTTGAAAACCCATTTAGATTTATGATTTCTAGCCCCCAAAATAAATTTTGTTCCTATATCCAATTCCTTCCCT
CACAGTTCTTTCACAATAGCTTCTTTTCTCCTGTAAAACCTACATAAACTCCAAAACATTTCTAGTTTTTGAAATCC
TAATCCAAGAGGTCACATCACTAACGCAACGTAGAAATCTTTGTACCAAAGGACAGAGGTGCAAGGGAAGTCGGGAGT
TGTGGCTGTGGGAAAGTGAGGAGGACTCTTTGGCATCTGGGCAGAGGTACAGGGAAGCCAGGATGGGGGCAGAAAGAAA
GTTTCTCAGTTTCTCTTGTGTCTACGGTTTCATGCTTAGGCCCTCTGCAGCAGCCCCAAGGCAGGTGAGGGTGATCAGCTG
TTCCAGTTTGCCTGGCACTGAGGGATTCTTGGGATGTGGGGCATTCAGTGCTAAAACTGGGGAAAGTCTTGGACAAATT
AGGACAAGTTGGTCAACCTACCTTTCCCTTGCTGCTGCTGCTGCTTTTAAAGTCCTCATTCCTTTGAAAATTGCAGTGAT
CTTCAGTTCTTCGTGATTTTAACTGTATACAAATATTCAAAGAGCTTGGACCAGCACATTTCTCAGAGAGAGCTCTACT
TACAGTAAATGTTATGTGACAGAGCAGGATAGCAAATAGATGTTACCTGGGGCAACTCTAATTGCTTGGTTCATGACTGT
CGGGAGTAAAGTGTTAATAGTTTCTTGCATATCCTCCAAGAGAGTATTACAAATACTATTCTGCACCAAGCTCTTTGTA
ATTTAATAGTGATCTTGGGGCCCTTTTGTATGGACTTATACAGATCATGATCTAGAGATAATGATCATTTCTTATTTGA

Fig. 9.310

CACATAGTATTGCATTATGTTATGTCTAAAAACCCATTTAACTCTATTGATAGTCTTGTGGGTCATTTCCAGTTTTTAC
TACAATTAATACAAATAACACTGCAGGCATCCTTGTGCCTGAACATCTTTGTGAGTTTACCCATTGAATAAAGTCTTAG
CTGTGAAACTGAATTTAAAATTTTCATAAATATTGCCAAGCTGTCTTCTAAAAAGAATATATTAAGTTACAATCCCACC
ATCATCACAAGAAATGCTTGTATTTCTATACACACTGACCCTGAGATTACAAAAACATATTTTCCCCATCTAATTTA
ATTGCAAAGAAAATTAGTCACCTTTCATACATTTATGGGCCAGTTCCCTCCTACCTCCAAAGACTCATTTGAATCAATTA
TCTTGAGGAGTTTATTTTATGGATATGTTAATGTTTTGAGTATATTAAGAAAATGAACTCTCCTCATTTGTGTAAAC
TTTTCAAAAAATTTTAATTTGTTTAGGGTTTTTTTTTTTTTATCATTTAGTCTTTTAAAATATTTGAGTAGTCAAATTC
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TAGATTCTCAATAAATGTTTTTTAAATAAATGGACAACGAATGCATGAATAAGTGAAGGATATAATTTATTCTTTTCAT
TCCTCCTCATGCTATGTAGAGATTTGCAAAATGAATGCCAAAACATATCTTTGAACAGAATTTTCATGGCCCAGTATCTT
CAGGCAGTGGTTTTCTGTTTTGTTCAAAATGATGTCCCTATCATCTGCATAGTGTGGCGTATGGGAGATATACGTAT
TGAATGAATACATAAATATAGAGAATAATGAGATAAATAGAACGTGGATTATTTGGAGTTCATCCTCCTGAGCTAATGA
TGGGCATTTCTGGAAGTGCTGGGTTGCAAATAAGTCCAGACCAGAGAGGATTGTGTTTTAAAGGCCCAGAATCTCAACTG
ATAATCCTCAGAAGCTCAGATTTACTTTTTTCAATCCAAGTAAATTAAGTGGAGTGGTAAGAAAAAGAAAAATGGTTA
TGACCATCAACCCCGAGAAACAGGGACCACATTTAATTGAATTTATAATGTCCCTAACATCTCTGCTGTGTTTGAAAGGT
TAAAAATTTCTAGAGAAAACAGCTTTTGCTACTACTTGCCAAAGTACTCATAAAATGGACTGATTGCTAAGAACAGAA
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CTACTCTCCTCAGACCACCAGCCTTCAGAGTAAGGGCACATCCTTCCAATAAGTGCATCCTTGTGAGGGACATTTGGT
GAGGTGCAACCCCTTCAATGTGTCCGTTTCCTGGGTCATTTGGGGTCTTAGGAAAATCTTCTTGTGGTGGCGACTGAA
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AGGAAGGATTGTAGGTTGGAGCCATACGCGACTAGAACTAAGACATGTATCCTTAAAAACCTCAAATCTGTGTTCTG
ATGGGAAAACCTGCTAAGTCAATGGTTCCAGACTTTGGAATTTCAAACAAAAAACTGTGAGGCCCAACATACAATT
ACCAACTCTTCATTTTTTGCCAAACAAAACCTATTTTCAACAGAACACACTAACATATACTGGAACCATGATTTTCATATTA

Fig. 9.311

Fig. 9.312

Fig. 9.313

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GTGCAATGGTGCCATCTCAGCTCACTGCAACCTCCGCTCCCGGGTTCAAGCAATTCTTCTGCCTCAGCCTCCCAAGTA
GCTGGGACTACAGGCGAGTGCCACCATGCCCGGCTAATTTTTGTATTTTAAGAAGAGATGGGGTTTCACCATATTGGCC
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GTGCCTGGCCGTATACCACATTTTCTTTATCCATTCATCTGTTGAGGGACCCTTAGGTTGATTCTCCATCTTTGCTATT
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GTAGTGAGATTGCCGGACATAGTGGTTAGTTCTATTTTTAGTTTTTTTGAGAAATCTCCATGCTGGTTTTTCATAGTGGCT
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GCATTTCTTTGATTATTAGTGATGTTGAGCATTTTTTTCATATTGGACATTTGTATGTCTTCTTTTGAGAAATGCCTATT
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TTGAGACCAGCCTGGGGAACAGACTCTACTAAGAATAAAAAAATTAGCCAAGCAGAGTAGTGCATGTCTGCAGTCCCAA
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CACTGGCCTATGCAGCAGAACAAAGACCCCATCTCAAAAAGCAATGACACACAGAAAAAGCAGAATTTTAATTTGTTCTG
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CTTGAATTGTGGTGCGTCTGCCATTGGTGTGGAAGCAGGGCAGAAAGAGGACAGAGGCATACCAGCAAGTACAGACATC
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CCTTTGGAAAGTTACCAAGGTAAGACATACCTGATACGTTCAAGAAACAATTCGAAGGTTCAATAAGGCCAGAAGTTAA
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AAATACTCCTCCCATGGCTGGTTCCAAGCTACCCTGGTTCAACAGTTCCTTTTGCAGAATTCCTTGAATCTTTAACACTT
AACAAGATCCAGTTCCAGCATATTCTGCAACAGCGATTCTGCAGTTCTCAATGTCAAATAATAGAACCAAAAAGTCAT
ATAAGTACTACAAAGAAGAAGAGAGAGTTGAGAGCCAGCTTAGCCAGGAAATGCTCCAGGGGGAAGTAAGGCTTGAGTT
GGGACTTATAGATAAGAAATGTAAATAAAAAGGAACAAACACTCCCACCGTGATGCACAAATATTGGATCATGTATGTG
TGTGACAAGTGTGGCCGTCTGTAGTGGAGGGTCCAGGTAAAATTTAGGCCATGAAAGCCTTGGTCTGTCAAACATAAC
ATTTTGTACATTAAAGTTAGTAGACAGCCATTAATGGTTTTTAAAAACAGATGGTCATAGCAATGTTTTGACAAAAGC

Fig. 9.314

AATTTTAAAGGAAGATTTTGT TTTTATAAAAAAGTTGAATTATTGATAGCAGTGAGACCC CAGAATAGAATAAGAGACCT
GCCTCATGTAAGTGAGAGTAATTCAATTCAGCAGTAATCTTGGAATTTGTAACTTCTGTGGTGAGAAATTTGTATCCTG
GGATGTTGGGTAAAGCCACTGTGAAGCCAAAGAAATTAGTGGCTAAACGTAGAGAAATTAAGATGAAGGAAGTGGCACA
TGGAGCCACTGTGTACTATAAAGTATTTTTATTGGTATTTAGTCTTGCTGTATTGTGTTGCTAATGATTGTATTGAATAA
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TTTTTAAAAACTTGCTTGAAACATGGAGACTTGGGAATGGGACGTCTATCATAGTAGCACATCTGAAATCCTTCTCATTC
CTGCTGTCATTTCTGTCTTTTCGCAGTCCACTTTGTCACCACCCCCACCATTCTTGTCTGCTGCAATCCAGCTTTC
CAGTCACTTTCTCTACTTCTTGCTGGCTGCAGCGTCATCATTTAAACAGCTCTGGGTGGTATCTCCTTTCACCAGAA
TTCCTGCAGCCGTGGAGTACGTGCAGAGCATCTTCTTTTGACCCTCTTAAATGCTGAGGGGTGTTTCCGACACACAT
TGCTTTGTAATTCAATAATAGGGCAGTTGTGGGTCTTTTTATTGCAGTTATGCTGATTTTTTAAAGCACTCTGAGTAA
GAAAGGAAAGGTAGTAAATATACGTGTACCTCGATGCTCATTGATCACGTTTGCACATGTTCTTGCCAAATGTTGTTT
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CTTGTTTACTCTGATTATATAATGGGGAATTGTATGCTTTATACAATTTTAATTTCTTGAATATATGTGCTCAAAAAC
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AACAAACCAAAATCATTGCAAAGTTATTTTTAAACTATTATTAATTATTTATGTAATCATAGATTATGGCTTGCAGAAGA
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TAACCTTTTCTCTATATGTACATTATGTGTTGATAAGCATGATATCTATTATACCTCAATTATTAATAACAATATACAT
CCTAACTATTAAGAACTTAGCCTGCAACCCTAATAGTGTCCCTTTCTTCTAGAACATAGATCAGCATAAGGAAGGAAG
CAAAGGAGGTCTCTAAGCACTCAAAATAGATGTTTACAATGTTCACTCACTGAAGATAATGTAGGTCATTAGCCAGAGA
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CCCAAATCATCAAAAATAACAAAATTATATTTTTGCTCAATCTGTATTGCTCTTCTATATTATGGTAAGATTTCTGTTTA
GTGCACTTAAAAGTAAGACTACCCCAAATTTAATCCCATCCCTTCAATGTCTCCTCCTGGACATTGCTCCATCTAT
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TCTGTTGTATCACTGGGTATAATGTCTGGCATTGTCAGGTGTGCATGCAAGGAAGGAAGGAAGTGAAGGAAGGTG
GAAGGAAGAGAACAAGGAGGGAAGAAGGGAAGGGTAGACACTTTGGGAGGCCAAGGCGGTTGGATCACCTGAGGTTGGG
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CTACTCTTTTTTAGAACATCTCGTATCTAATACATTAGAACTCCTGAAGCCTCTGCTCTCAAAATATACCCAGAGCCCAG
CATCCTTGTCACCATCTCCATGGTTACCATCCTATTGCAAGCCACCATCTTCTCTGGCTTGGATGATTGCAGTGGCTGT

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CTAATAGGAGACCCTGCATCTCCTCTTGCTGCATAACAGTCTATTCAAAAGACAGCAGTCAGTGCAAATCATGTCATTC
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TAGGGCCAATTCCATTGCTTTAAGTCTTTGCTTGTGTTTCATCTCCATTAGGCCATCCATGACCATCCCATTTTTTAAA
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GCCTATTTTGTTCAGTACAGATATATATACACAACAGTGCCTCAGCCATGCTAGGTTGCTCAGTACATTCTTGAAT
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GTGGGATCCTAAATAGCTTTCTATGATCTCCCTAGAAAAGTGTAGAAATTTCCCAGAGAATAAGCCAGCATTTTGTGTA
CCATTCTGCAATTCCAGGATCATGCTTGCATAGTCATAGCTTGGAAAGGAGGCAAATTGAAACAAGTTGAAAATCTGCAG
GAACTATCCCAGTGAGACCACAGAAAAGCCAGAAAGAAAGGTGGGATTTGGGGTAGAGAACGAGCCACTTCTCAC
TAGTTTGCATGAAGCATTGAATATCCCAAGGGAGAAAACATTGAAGTTCTATGAGACACCAAGAAAAGTGTATAGATTA
TCACAGCATATGGATAGAGGTCATTTGTTCCCTCCCTCCCTTTAAAGAAAAGTAGCTTTGCAAGCCACAGTTTAGGAAC
AAATGGACTTTTTTGACATAGTTCAAATCCTCCTGGGTGTGGAGGCATTGACAGGAGCAATGTCATAAATTGGTTAAAGG
TTGCAGTCTGGAATCAGGCTGCCCTGAATCCCAGCCCTGCCATTTACTAGCTGTGAGACCTTGTGCTTCCCTAACCTCAG
TTTTCTTTTCTTTTCTTTTCTTTTGTAGACTGGGTCTAGCTCTGATGCCAGTCTGGGGTGTAGTGGCACCATCTCAGCTCAC
TGCAGCCTCTGCCTCCTGGACTCAAGCGATCTTCCCACCTCTGCCTCTCAAGTAGCTGGGACTACAGGCATGCACCACC
ATGCCTGGCTAATTTTTGTATCTTTTGTAAAGATCGGGTCTCGCCATGACACTTAGGCTAGTCTCAAACCTCCTAGGCTC
AAGTGATCCACCTGCTTCAGCCTCCCAAATGCTGGGATTACAGGCATGAGCCACGGTGCTCAACTAACCTCAGTTTTC
ATAATGGTAAATAGGAATACCGATAGCACCTCCCTTGGTATAAGGATTAAATAAGATAATCTACATAGTGCTTGAAC
AATGCCTAGGCAGTGTTTCATCAACAGTTAGCTGTGTGCTGTGTACATGCTGGAGCTGCTACCTTGTAGAAATTTTAAAT
TTAAATGGTGTGTAGTTGGAGGTTAGGAGTTACTAGGGGTATCGCCTAGGGAAAGACAGTTGGTTACCCAGACAGGTC
CTTACAGAGTAGTGTTTTCCCTGGAGAATTAACATATATCCAGAATCTGTCTCAACCAAGCAGTCTCAGAAAGGTGAT
CTTGTCACAGCCTCTGAGTAAGCTGATTCAAACCTCTCAAAGCTCACAAAGAGCCTTAAAGCCAGAGTATCAGCTGATTC
CTTAAAGCTACAAAGTGTTTTGGCCTTGCCAACATACGCATTTCTCCCTCTTATGGGTAGGTTTAGAATGCTAAATAGT
ACATAACATGGATTTAGAATAGACAGATTTACATATGATCAGAAGGCTCAGTGGTCATAGTTTTTGAGGGCCTAGGACAT
GCATGTGAGCAGTAGGGAGATCATCACAGGAGGATGGGCACACTCTCGAGGCTCATAAGGCGCTCCCTCTCTCCATGG
GCAGCACAAGGCAAGTCTCCTTTGTCTTTGATTTACTTGCCAACCACCCTCATTCTAATGGGAACCTTTCTTTTCCAAA
TATGTTAACACCTTCAAGTAATTACTGTCAGCGTCCAGCCTGGGAGTCTCTTGACACAGTTTTTTCATAACTCTAGATG
GATCTCATAAATTTGCCCTGAAAGATGGTAAATAATTTTACTTCTCTTACTTCTGGACTCTAAGGGGGATACTTCACTC
TTGCTCCTTGACCTCTCCTCTCTAGATCCCACAAGCCAAAATTTGTTGGAAAGGATTAATATGAGCTGGGCCCCAATAGG
CTACAGCATGCACCAGCCCCCTAAAGCCAGCTTTAGGATTGGGCGGAGTGTGAGCCAGCTTCCACCCCACTTGGCTTAT
TCAGCGACCTCCTGCAATTGTCCTCTGCTAGCCCTTGGCAGATCAGAAATGTTCTAAAGATTGACCTCTATTACTTTGG
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ATAGCACCCATACTTTTCTTCTCAGCATTTATTATGAATGTAATCAAATCATACTTTGTGCATGTTTTGTTGTGGTTG
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AACTTCTATGTGCATTATGAACTGATATAGTGGGTCACAACCTATCAGTAAGAAAAAAGAACTGGAATAGAGTAGAAA
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TCTGCCTCCCTGGTTCAAGTGATTCTCCTGCCTCAGACTCCCGAGTAGCTGAGATTACAGGCACCTGCCACCATGCCTG
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CTCAAGTGATCGGCCCCGCTTGGCCTCTCAAATGCTGGGACTACGGGCGTGAGCCACCGCACCTGGCCAATATTTTTT
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TAATCCTTTTCTTAATCTTCCCTATGACCAATATTACCATGCACACATGCACACACATACACACATGTGTGCACATAC
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GGATACAAAATAATAAATTTCAAATAAGTCTTGACTAGTGAGTGTAGAAGGACAGTGCTACATCTGAATCTCCAGTTG
GGTTTACGCATTTTGTATTTATCAAAGGCAGTAGTGATTAAATGTTGATAAACACTCAGTTTTCAGTTTTCTATTGC
TGCAAAACAACTATTCCAGAAAGTGATGTCTTGAAACACAACCATTTCTATTTTGCAGTCTGTGCTGGGCTCAGCCGGGC
AGTCTGTGGTCTTACCCATGGTCCCCAATATGGCTGCAGTCACTCCAGTAGACTGGAGGCTGGGCTTAGCTGAGATG
ACCTGCTTTCTTTCTTTCTTTCATGTAGTCTCATGGCTTTTCTCTCTCTCCATGTGGATTCTCATGTGAACCTCTCCAGC
AAGGTAGTCAGAAATTTCTTCCCTGGCAACTCAGGGCTGCCAAGAACACAAAAGCCAGGCTTTCTTAATGTTTAAGCCCA
GAACTGGCATGTGGCCAATTTTTTGTGTTCTGTTGGTTAAAGCAAACCGAGTCAAGAAGAGCCTACACAAGTAGTAATA
TTACAAGTATGATGAAATATCATTTCTTCTTAGACCAAACAGCATATTCCTCCCTCCACTCTTACCCCTCTATAATGTT

Fig. 9.316

TTGCTTTTGGGAGTTCCTTTATATGAAATGCCTCCCCTCTACTTCTCCTTCTACTGAAATCCTACTCATCCTGAATCCA
GCTCAGAGCTAACTCAGTGATTTCTTCATACATCTCTATTATCTCATCCTGGAAATAATCACTCCTGCCAGGTGCAGT
GGCTCATGCCTGTAATCCCAGCACTTTGGGAGGCTGAGACAGGCAGATCACTGAAGGTTGGGAGTTCGAGACCAGCCTG
GCCAACATGATGAAACCCCGTCTCTACTAAAAATGCAAAAATTAGCCAGACATGGTGGTGAATGCCTGTAATCTTAGTT
ACTTGGCAGGCTGAGGCAGGAGAATCACTTGCACCTGGGAGGCGGATGTTGCGAGCCAAGATCGTGCCACTGCACTCCA
GCCTGGATGACAGAGCAAGACTCCAGCTCAAACAAAAAAGAAAAAAGAAAAAAGAAAAATGCCCTCAGCACTCTCATA
GCATTATGTCTATATTTTCATATTATGTCTTGACATATAACTTTACACACACACACAAGCACACACACACACAGATGTAG
CTTGCTTTTAGATTTTAAAGTTGCTTATCTAAGATTGGGTGATCAATAACTGTTTTTTTCAAAAAACATTAAGTGGTTTT
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TATTGGAGAAAAAAGCTGATGTATGCTACCAGCATAAAACAACAATTGCCCTTTACATCTTCAGAAACCCCTGTACTG
TGCAGTCATTTTCAAACCTGGATTTTTGTCTTGTGTTTTTTTCCAAAGTAGTATATATTTAACTTTGTAAATGGTGTGG
TGTTTTTTGGTTGTTGTTGTTTTTGAGATGGAGTTTCATTCTTGTACCCAGGCTGGAGTGCAATGGCGCGATCTCAGC
TCACCGCAAACCTCCGCTCCCGGGTTCAAGCGATTCTCCTTCTCAGCCTCCTGAGTAGCTGGGATTACAGTCATGCGC
CACCACGCCCGGCTGATTTTGCATTTTGTAGCAGAGGCAGGGTTTACCATGTTGGTCAGGCTGGTCTTGAACCTCCTGAC
CTCATGATCCAACCTGCCTCGGCTCCCAAAGTGCTGGGATTACAAGCATGAGTCACCACGCCCGGCTAACATGGTGGT
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TCAATTAGTCTTAAACTACAAGTTGTTCTCTATACCAGCTACACATGTGCATTGGTAGTCTTAAATTGTGATATGAG
CCCTTAGGTAGTTTACACCTTTCCAAAGGAAGCTCAGTGAACCTCCTGAACTGTGTAAAGAATTGTGTGTGCAGTATG
TGCATGTACATTATCTGTATTTTTCTGGGTGGATGATCCAGTATCCTCATCAGAGTCTTAAGTTTGAGAAACCCAATTA
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TTTGATTTCTATTTTCATATCATCTGACAGTTTTGATTAAATCCTTTAACCTCAGGATTTGTAAGGGGTAAGGGGAGTG
GGTTAACTAACTGATCTGATTATCAAATAATGCTACCATTGAGGACTTATGCATAGTTTCTGTGATTCTCAATAGAATT
TGATGGTACCTACAACCTCTACTCCTTGTTTTAGTGTCTATGTTTTTCCACTGTGCTTAAAAACATTTAAAGGGAGAGG
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GGAAAATTCAGTTTACAAAACCTCTGAAAAGTTCTAAGGCAGAATTTTGCTGCCCGGTTTCTCATAATAGACCCAGC
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GTAATTTGGGTGTACAAAATGCCTGTAGTTCAAAGTGCTTTACTCCTCTGCAGTGGCAAGCTGAGCTTCTGTTGGCTG
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TCAAATGCCATAGGCCTTAGCTCCAGGACATTTTCTGCCTGTCTCCTTCCCTCCTTCTTTCTTTTCTTCTCTCTCC
TTCCCTCTTTCTTTGTTCTTCTTTCTTCTCCCTCCCTTCCCTCCTTTTCTTCTCTAGTTCCCCCTTTCTTTCTTTCTT
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TGTTGTGAATTACATGGGGCTGTGGTAAATGTGGCACATTTCAAGGCTATGTATCCCTTTAGATTCTGGTTCAGTAAG
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CAGTAATCACACATGTAAGGGCTTTGGAGTAAGATGGACCTTGGTTATCCAACGCTTACTGTGTGACTTTGGTAAATTA
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GACAATGTATTTAAAGCACCCAGCATACTATCTGGTTCATAGTTTACAGTCAATAAATGTTAATTCCATTATTTGATCA
CAATGTAAGACTATTGAGGTTTAAATTTTGATTGTTTTTTTCAAAAGTTTGTAAATCTGTTAGGAACAATTTCTAAAGTAAA
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CAATATTTAATGCGAACTTCCCATTTTTGAACAACGTGTGTGGGCCAAACAAGTCACATCTGTGGACCAGATCTGGCC
TGAGGATTGCCAGTTTGCAATCTTGACCTAATTGCCCTTAATTCTTCACTTCTCTTGACCTGGTAAATACTCCATTTAA
ATGAGTATTAGGTATGTTCTGGGCCCTTTTGGACTGGGTCTTTGTAAATATATACAAATTTCAAAGACTGACCTTTAGTT
TTTCAGGTTTCAGAATTGATTTTCTCAGAACCCAATTAGATCAGGTGCTGTCACTGACACTCAGAACTCATGAACTTTAT
GCAAGAGCAAGAACATGAATTAGGGAATTCACAGTGAGAAATATATTGACAAGTAGACAGGATACCATATTGGCCAGAT
CAGTTCATTTCTGTCTTGTAGTTTCAAGGCTTTGAGAAACAACTTCTAATTAGAAGCTGCTTTGACATATATGGAGTCA
GAACTAGGAACTGTAGACTTTAAAGGGCTTTCTGCAATTGGGTTTTGAGATTCTTCTCCTTTTATGGTAGTCTTAAC
ATTCGGCTGTTAAGTGTATGCTCCCCCTCCCCAGGCATGCCAAGTATATACTGACCAAGTGTCTTCTTTATTTAGCTA

Fig. 9.317

CCATATGAGTGGTTCTCAAACCCATGCCCTTGTTAAAACACAGATTACAGAGCCCTATGCCCCACAAATTCTGATTCAG
TACTTCTGGGGTGGGTCTGAGAATTTTCCTTCTTATAAGTTCCCAAGTGATGCTGATGCTGATGAAGTCTGGATGGG
GAACCCCTTTGAGAAGCACTGGGTTCCTCTATCATCTTCAGCTGACAGGCTTTTTCCCTTTGAAGGGTTACCGCTAT
TGTCTGTGTCTCTGCTTTAATATGCTAATATAGCATTATGGTCATATCCAGATCCTGAGGTTGGAACCTGGGTCTTAGA
ACAATTTTTTAAATTGTTTATATTTTTTAAAGGTATTTGATTTATTAAAACTATTTGATTACTTTATTTTCATCAAGTCCAA
GGTGCCATTGATTATAAAAAACATGTCTGGATTTTAGAGGCATTAAAAATGTAGGGGCCAGGCGCGATGGCTCATGCCCTG
TAACCCCAAGCACTTTGGGAGGCCGAGGCGTGAGATCACCTGAGGTCGGGAGTTTGAGACCAGCCTGACCAACATGGAG
AAACCCCATCTCTACTAAAAATAAAAAATTAGCCAGGAGTAGTGGCACATGCCTGTAATCCAGCTACTTGGGAGGCTG
AGGCAGGAGAATCGCTTGAACCCAAGAGGCGGAGGTTGTGGTAAGCCGAGATCATACCATTGCACTCCAGCCTGGGCAA
CAAGAGCAAAAGTCTGTCTCAAAAAAAAAAAAAAAAAAAAAAGTAGGGAAAAAATTAACAGTTTCAGAGGTACTGCAA
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GCACTTAAAAAAAAAAAAAAAAAGTTGATTATTAGACTCACTGTTTTCCTTGCCAGTCAGGAAAAGATATTTAGGACAAGAG
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AAGTTTAGAAGCAGACTCTGCTTTAGACTGAATAATCCCTGAGGTTCTTGGGTTATTTGAAAGAGGGGTAGTTTTCAA
AAAGAGAGATATTAGATTTCTATTGAAAGGGCAGCCCTGGTCTCCAGTGATTAAGTGGAAAAACAAAAGAGATATAAC
AATTTTTTACATCTAAGTACTGTGCTAGCTTCTGTGGATCTAGAATCAAATGAGACAAGATGTAGTATGACAAGCAGTTA
GACTCTCAAGAAAATCATTCAAGTCTGTTTAGGAAAACCTGACATTCATTCAATTCAGCAAACTATTGAGTAGCTCCTGTG
CATCAGAACCTGTATTACAGTCAACTGAAATAAACACAATTTCTACACGAGTGGAGTTTTCCAAATAGACTAAGATGT
CATAATGGACTGTGTGTAGAGTAATGTTCAAGTTTTTTTTTTGTTTAAGCTTCCCCATCCCCCAGAAAATACTGGATATAT
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GGAATGTTCTAAGCATGACATTGTGAAAGAATGCTAGAATCTTTTTTCATATGAGGGGTGTTATGGAAAATAAAGAAAGT
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AAGTAAGTGATGGCCATAATCGAAGAAGAGATAAGAGGTAAAGAGCAATGGCTTAAAGCAAAAAAGCCTCAAAACATTC
AACTTTCCTTTGTATAATAGTGGATGTTATTTTGAAGAATGTCAGTTTCAGGAGATACCATAATCATGTGTTTGTCTGTA
TTTTAAAAGCCACCACCATAAAAGATCTAGAGTCACTCATGAAGTTCAAGTACCAATTTTTTACCCATGAGTGTGGAACA
TTCTGCTCTTTTACAAACAGTAACCTCGTACTGTACATTTTGCGGCGCATCTCTCATTTTTGTTGGTTATTTAGTGGCCA
TGTAACATGTACATGACTTGTGGTGAATATGGTGATTCTCACTTTATAACCAAGAGGGTGGATGTTACAGCATATGA
GCAGTTATGACTGTAAAGCCTGAAGTGTGAGTCACAGGGTCTGACCCAGGTAGTAAGATGTGTTATTTGTTTCATGTTGG
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GAATTCCTGCACCTAGTTCTTTGGCTAACCAGTTGTGTCTTTTGGGGGAAATTTCTTAATCTTTCTGCTCTCCATTTCCC
TTTCTGTAAAGTAAGGGATTAGACCAAAATTCATCCAGGATTGAGAAATCTATGGTGTGACAAAGACTGCTGGTGTCTGA
CTGAATATCCTTACAGATTTGTTTCATTTAGTAACCAATATAATAAAGATGACTGACTGAATCTTAATTATATTGGGTGA
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TCCTGCCTAGAAGCACTCACTTTTACTCCTCCTCCCGTTTTGTGTAGTAGTGATTTGGTGTCTGGTGTCTGGCTGGAGCATC
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GAACCTAGGTCCCTGATAAACATGGTGTACCAATTCGAACCTTGACCTGTTTCTAGATTATTTTCTTTACTTTTCTTTT
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CTTACTGCAGCCTTAGCCTCCTGCACTCAAGCAATTTCTCCCACTTCAGCCTCACGAGTAGCTGGAACCACAGGTGCATG
CCACCATAACCAGCAATTTTTTTTTTTTTTTTAAATTTTTTGTAGAGATGAGGTCTCACTATGTTACCCAGGCTGGTCTCAA
CTCATGTACTCAAGTGATCCTCTTGCTTGGCTTCCCAAAGTGCTGGGATTAAAGGCATGAGCCAACATGCCCAGGCTA
CATTTTCTTAATATGAGATAAAAAATAAACCTCTTTCTTATTGAAGCCATTACTAGATGCCTAATTCATTTCTAACTAA
CATATTAGCATCAGATTATCTTTATGTAATTTCCATTGCTAGGTTTTCTCTTTGCAGTATTGGAGACAATAGCTTACCA
ACTAGCTTGGGAACCTTCTAGTGCTATTAGAGTTTCAACACAATTTACCAAATTTCTAAAATTTATTAGTTATTGGATA
TATGAAAACATAATCACCATATGTGAAGAAAAAACCAATGTTTAGTACAAAAATTGGGAGGGGGGAATATTATATTA
GAGAAAGTTTATTAATCCAGAAAACCACAGAATTTAAAAAATATTGGAAGTTGGGAATTTGGGAGTTAAAGGTACATTT

Fig. 9.318

GATCTTTGGTTAATGGCAAGTTCACATTTTATAATAGGATGTAAAACCTATTTCTCACTGCACCTGTACTTTGATCTT
CTTGTCTGTCATTGCAATGTGATTATGAGGATTACTGGGTCACCTCAGCCTAAGATGTTTTTGTGACTTAATTTTCTA
GGAGAGGAAGGGGATCATTAAAGGATATCATTAAAAGGAGAGACACAAGGCTTGTAATTGACCTTCCTTGAAAGTAGTC
CCATCTGGAAGATTTTATCAGTAAACATTTATCAGTGTATTTGGCTTTTAAGCAGACTCTTCTCCTCTTGTCTTTGAAA
CAAACAGGCAAAGGCTAAAAAGGAACACGTTAGTGCTCAAAGTTTGCTTCTTGTGCATGCTGAATGGGAAGAAAAATAA
TATTTAAGAAAAAGCTTTCTTCTAAGTTAATTACATGATTCTTTTTTTCAGTTTTGCTTCAATGTTTCTGTTTGCCAAA
TTTTAATTGGCAAGTTATGGTGTCTTCAATGATTATGCCTTGGTGTGATTTGCTTGGATGTGTGTTTAGAGAGTGGAG
GGTAGAAGGGATGAGATTTTTGTGAAATACAGTGAGCAGGGGCTTAGAAAACCTACTGCAGTTCTCTGTGTGACTAAGC
CCAGCACTAGTCTGAACTTCGTATTGTACTTCTCTGGAATGCAATAACTATCGTCAGGAAGACAAACGTTGCTGTGGC
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TCACAATCCTTGTCAATTTGCCAATTCATATAAACTTTCAGAAAGTGAAAATTGCTTTTAAGAAATTTATTTTGAAGTT
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CCATGAGGTTTAGGTAGACCATAGATACTGTTTCAATTAGAACACTTCATGAGGATTTAGTTTCTCTGTCTATGGGTCC
TGTATCTGAACTCTCCATTTTTTATTCTTTTGTTTTGTTTTTTGTTTTTCTCCCTGATGCCTGATATCAGAAGAACT
CTCCATTTTTTGGCTTCATTCTCTGCTCTATCCACATGAGACAGACCAATGTCTGCAATCTAGTGCCATACCTTTCCAGG
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CCAATCAGTCAATCACAGTGGTCCCATCTGACCATGGGGATGTTCCCCAAAGGAAATCAGGGTGTATTTTTTCAAAGG
AAGAAGAAATAAATGCTGGACCTTCATAACAACCAATTGCCCACTAAAAGTGCAGTCATATTTCTTGATGAAATAGAAC
CTCTCCTTGTATACTGGGCTCCATGACACATTGATCATGGTTAGCTAGAAGATTCTGAAGTCTTATTATCCACTTTGCC
AGTTACATTTATTTGTAGCTCAAGGGGACTGTTAAGGGAAAGGGGAGCAGTGCATAAAACAAATCCCATTCCAATATTA
TTAAAAAACCATTGCTCTGCCCTTCTTTATTACAACTTAGTTCCTCATCCTCATCTTCCCCAGCTCCTCCTTTTCA
GCTATTTCTCTGTAGGTGCCCCTACCACCTAACTCCCACAGATTCCCCCATCTCCTTTGAAGAAAAAAGCTAGATGCTC
CCACTCCCTTTATTTCTTCCACAGTAACCAGCTTAAGTACACACGGCTTTGTACATGGTACACAGTTAAGTAATATTT
GTGAAATAAATACGGAACACTTAAGGGAAATAAAAAGCAATCTATCCTCATCAAGAATGAGCTGAGAATCCCTGAGATA
GCCTATTTAGCAGCTTAGCCTGAATTCTGTTTTAGTTTTCTGTTTTCTGGAACTCTTCTTATTTTAAGAAAGAATGATTC
TTATAATTATTTTCATATTTGTTGATATTATCCAGTCTTAGCCAGTTATCAAAATGGTCTTGAGAATTAGGAGGGGAAAG
CATAATGTTATAAACATTTTCAGCATATTGTGCAAGTAAAGTCCCTGGTAGTCTGGGTAATTCTAGAACTCGAGTAGACTT
GAAGTGATATCAGAGGAAAGCTATTACCCAAGTTTACTGCTTGCTAATCAGGCAGCCAGCAATAATGAGATGGTACAAT
AGAAATGTTCAAAGTAAGGTTTCTTGGCATAACATAGCTATTTAATTATTTTATTTCTGAAAGAACTATAGCTCCACTGT
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TATCCTTTGTAAAAGAATACTATAATACCAAGAAAGGCTTTTGTCTTCATCTTTTGTAGTTTTTGGTTATTTTTTGTTC
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AGCCTAAAAATAAGAAGATACTCCTCTACTTTGTATACAAGATCGTTGAATGGTGGGTCAATTAGTTGATAGCAAAAAG
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GGTGTAGTGAAAAAGAAGTCTCAGGAGCTCCAGAGTCTCAGAAGAATGACAAGAAGACCCCTAATTCTTGCTGTCTC
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AGGCAGCCCTTTGTTTAAACCTTTTTTTTGTCTCAGGCATGGATTAACTGCAATTGGATTCTATGTTAAAAATGTATTTT
TTTGAACATTTTGTAGGCATCACTCCAGTATTTTCTATTGGTGAAATCCTGATCTTTTTTCTGTGTAGAAATTTTTTGTGA
CCTTCTTTTCTTTACTGAAGTTCCACATTTTCATGATCAATTGTCTTGGTCTGCATTTTTATCCATTGCCCTGGGCACG
CAATAGACCTTTTCACTCCAGAGATATGAGTCCTTCAGTTCTGAAATGTTTGTCTTATTTCATTGATAATTTTATGCC
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TTGTTTTCTTCTTTTTTGTCTTTTTTGTTTTTTGGTCTTCTAGAAGATTTGTTTACACACATTTTTATTTTCCAACCTT
ACTATTAATGTTTATTTTGGTCATGATTTAAAAATAATTTAGCACTCTCATTCTCTCCTTTTTTTTTTTTTTTTTT
TTTTTTTTTTGAGACGGAGTCTTGCTCTGTCAACAGGCTGGAGTGCAGTGGCATAATCTTGGCTCACTGCAACCTCTGC
CTCCAGGTTCAAGCAATCTCCTGCCTCAGCCTCCCGAGTAACTGGGACTACAGGAGCATGCCACCACACCCAGCTAA
TTTTTGTATTTTTTAGTAGAGATGGGGTTTACCATGTTGGCCAGGGTGATCTCGATCTCTTGACCTCATGATCCGCCCA
CCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCAACATGCCCAGCCTCATCCTCTCCTTTTTTAATGGTATATG
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AGAAACAATGTCAGAAGTTCTGTGATCATGGATAGGGCTCGTCAACTGTAGGGTTGCACTGTTGCATCATAGGTTGTTTC
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GCGCCTGGGGCATATAACTGACTGCTAATGTCTGGGGAGCATGACAAATAAAAAAGTTGGGTTTCTTATTGCAAACTCT
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AATCTCTCTAGCCGATTTCTCCTCCAAAAAAGAAAGACTCCGCTTCCAGTAGGTAAGGGAAAGTAGTTAGTTTGGATCCAAA
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GTACTTTTACAAAAATCATTTTTTCATGTCCATTATTTAGGAGTCTCTCTACTCTATCAAAAGTATTAATTGACCTATTT
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ATAATTCATCTTTTAGGCTTTGTAAAATTATCATTTTATAAATTTTTAAATTGTGAAATATAACAAAATTTAAGAAAGT
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TCATCCATTAATTGCTTGCCATATGCTTATCACTGGAATATATTTTGCAGCATCTCCTGACATCACTATTTATCCCTTT
TACCAAAAAAAAAAAAAAAAAAAAAAGAAAAAGAAAAAAAAAACATGGCTACTGATAACTTGGGAGCACTCAACGAAGG
TCCCGTCTGAGACTCTGTAGAATATATTATAGAATGCAAATATTCTTTGGATTCCGTCTTGTTGTTGTTGTTGCTGTTT
CCTACCCTACAACCTTTTGGCATCTCTACCCAACATTTTATAATACAGGACTGTCTCTGAGGTTTGTTTTCTTTATGGT
TGTTCTCATGTATAAAAGAGATGATGATAGCGTTTCTCATGCCAGTATGTCGTGTATGATTTATGTTGTAAACAGTGC
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TCTATATCACGGGAGGAAACATATCTCCCTGTGGAGAGATGTCCACTGCTTTATCGACAAGGCACAAAGCTGCGAACAG
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GGTGTATTCATCAACTGATCACTATTAGCMACTGGCTAGAACATGAAGTGTGTTTGTGTTTTTCAAAGAAAGATATGA
TGAACTTTATATGCTTTTACACAGTTCTGATATTTTAAACAATTTGACTAATTTTATAGTTTTATTCCTTCCAGAAAT
TCCTTTAACTGTGCTTATCCCGTAAGTAATTGCTAATGTTCTTAAACTAATCGAGAAAATCATTTCTATTAGTCCCTAA
ATACCCAGACTTCATACCTTCTTGCTTCCCACTCTCCTCATATCTAATCTCTCCCTTAGGTTTAGCAACAAAATGTGCA
GCTTGACRTGGTCAGTGGGTGTCTGAACATTTAGTGCTAGTGTTTCTCTCATCTCTCTGCTTTTATTTGGACTTCTCA
ACTCTTAAAGTAATTTTAAACATATTATAAGAAATCTTTGAATTTATAACACTTTACAGTTTATGAAAGACTATATAAA
GGTTCTTCTCACAGGCTGTAAAGGGGTACAAGGACAATGAAACTCAGATGACATCCTTAAATGCACTCTACTATTATAT
GGCTAATTTAAGATTAAAAATCTGGTTTCATAATTATTAGTCTAGTGGTCTTTTATAGGCAGCATAATACAGTTGTAA
GGAGTACAGACTGCCTGTTTTTGAATCACTGGTCCATGCTAACTAGCTGGCTTACCCTTGGGCAAGTTACTTGAGGTTC
TCTGTGCTCAGTTTCCCCAGCTATCAAATAGATCTAAGAGTTGTGAAAATTCAATGAGTTGATACATATAAAAAATACT
TAAAGCAGTGCCTGTCCAACATAAATGCTTAATAAATGTTAGCCCTATTATCTTTCTCATTATTTAGTGTTAAAGAGC
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AAAATTGATACCAATTAGTTTCTTTATCTTGAATGAAATCAGTATTGTAGTCAGAGCCAAGCTACTGACCTGGATTAC
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GGTTTAAGAATTATGAGGAGGAGAGAAGATATTGTAATGAATCCCTACTATTCTGTTTATTTTTTATTTCTTTTAACT
TTTTAAGTTCAGGGGTACATGTACAGATTTGTTACATAGGTAACTTGTTGTCATGGGGTTTGTAGTACAGATTATTTCA
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GGTCACAGTGTCTGTTGTTTCCCTCTATGTGTCCATGAGTTGTTCATCATTTAGCTCCCACTTACAAGTGAGAACATGCA
GTTCTGTGTTAGTTTGTCTAAGGATATTGGCCTCCAGCTCCATCAGTGTCTGGAAGGACACTATCTCATTTCTTTT
TATGGCTTCATAGTATTCCATGGTGTGTTATGTACCACATTTGCTTTATCCTGTCTACTACTGTTGGGCATTTAGGTTGA
TTCTATGTCTTTGCTATTGTTAATGCTGCTGTAATGAACATACACATGCATGTGTCTTTATAATAGAATGATTTATATT
CCTTTGGGCATATAACCCAGAAATGAGATTGCTGGGTGTAATGCTAGTTCTGTTTTTAGGTCTTTGAGGAATTGCCACAC

Fig. 9.320

Fig. 9.321

TGCAATTCAAGTAATATTCATTCTTGAATATTGAATTAAGCAAATATCTTAGAATATAGAGCAAAAAGCCAAGGATATA
AAATATGATGGAAAAGTAAAAAAGATATGAAGAATGTGGTAGAAGTGGAGGAATGGGAAAGGATGTGGGAAAATAACAG
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TCAAGCAAAACAAAGCAACGGGAATCCTATACAGAAAACCTAGGATACAAAGAGCATTGGAACCAACCCAGATGTTTAT
GAAAATAAATCCCAGTATGACAACCTGTGTAGTAGTAGGCTTAGAAAGCAGGCAGCCTAAGTTAGAGTCATCAGAAGGTT
CCAGGAAAAATGCTTTCAAGAAGAAAGTAGACATCACATTTTGAATTGCATAATTAAGAGCCTTTTAAACATTTTAGGGC
AAAAGTAACTGCATTTAACTCATAGATTAATCTAAGACTTGATATCTTTATAATATTGACTCTTCCCATGTAGGAACT
AGATATATTCTTCAAGTTTTTAAAAATGTCCTTCAGTGATGTTTTGCAATAATATTTATACAAGAGTTTTACATTTTTC
GTTGTTTTATTTATACATATTTGATATGTTTTATTACTATTATGAATAGGATCATTTATCTATTTTCTAACTGGTATATA
GAAAAAACCTGTTGTGGATACATGCTTTATAATGTTTGACCTTTCTTATAAATCATAATGGTCTACCTTTTTTTATACC
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TGCTGCTGTGAAGAAATACCCTAGACTGGGTAATTTATAAAGAAAAGAGGTTTAATTGACTCACAGTTCTGCATGGCTG
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GTTTTTGAGTAGCGAGTGGCAAGAAGCAATATGACGATGTTGGTAGGTAGAGAGATACTTCTTCACCATCTGGTGAGCA
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TGATCAATTTAGCGTTGAGGAACTGAGGGCTACAGAGTGCCAGGGTCACGTTTCATCACACTTATTTGTGTTTTCAAAGC
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CACAGACTGGGATTGGGAACATGGGAAACGTGTGTGTGTACAGACACAGCAACCTTTTCTATGTGTCTATGTTTCAGC
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TCTCCCCCTCCTCTCCCCCTCCTTTCTTTCTTTCTTTCTTTGACACTGGGTAGAGTGTAATGCAATCATAGCTTACTGCAGCC
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GCTTATT
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ACCATTGGTTTTGTGAAATTATGCATGCTTTTTTCTTAATTTTCTCTAATTTTTTCTCTATTTTCTTAAGATTTCTACCATGA
GATTTTTTCTCTTAATGTCTAGTTTTTATTACATAGGTAATAGATAAATATATTTCATGTCATAAAATATTTTTTAAATTC
GAATATGAGAAAGTCTTTTTTCATGCTCCTTAGTCTTGCTGTCTTTCTTGAAAGTATCCACTTAGTGATGGATACAATA
TGGTGTATATCTGCCTTAATCCATTTTGTGCTGCTCTAACAGAATACTTGAGACTAGGTAATTTATCATGAGCTGAAAT
TTATTGGCTCACAGTTCTGGAGGCTGGGAAATCTAAATCAAGGTGCTGGCATCTGTTAAGGGTCTTTTGTGTCATCA
CCACATGGCAGAAAGCAAAAGGGAGAGAGATAGACAGATTGGGAGAGAGAGAGGAAAGGGGACCAACGTCTCACTCTT
TTATAACAAACCTGCTCCAGTGATAACAGGATTAACCCTTACATCAGGGCAAAGCTCACATGACCTAACACCTCTTAGA
AGTCCCAACTCTTGCTGGATGTGGTGGCTCACACCTGTAATCTCAGCACTTTGGGAGGCTGAGGTGGGTGGATCACTTG

Fig. 9.322

AGGTAAGGAGTTCGAGACCAACCTGCCTGAGCAACATGAAGAAACCCTATCAATACAAAAAATACAACAAAATAGCTG
GGCATGGTGGCATGTGCCTGTAGTTCCAGCTACTCTGGAAGCTGAGGTGGGAGAATCACCTGAGCTTGGGAAGCTGAGG
CCACAGTGAGCTATGATCATGCCACTGCACTCTAACCTGGGTGTTGAAGTGAGACCCTGTCACCAAAAAAAAAAAAAA
AGAGTCCTACCTCTTAATACTGTTACAATGGCAATTAACTTCAGCCTGAGTTTTGGATGGAACAAACATTCAAACCAT
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CCTTTTTTAAATATTGCAAACCATACTACACTAAACATTGTCGTGCATACTTCCTTGTGTATTACATTGGTTTTTAAATC
AATGAAC TAGTTGTAAACATGGAGAGAAACAACAAGGAGGGAGATGGATATAAGAGAAGAAGAGATTAAAGGGGATGGA
ATCAGTTGTTTCTGGAGCACAGAATATTCACCAAATTTGACCAGATGCGCGGTGCTCATTTAGGAAAGAGCAACTGCA
CTGGGATGAAGAGGTTTTTTTAGGAAATCAGAGGTCTTCAGAGAAGTTTTGGTTAAATCTGCAGTATACTACTACCAAAA
TGGTTTTTGTCTTTGTTGTTCTTATGCAAGAAAGACTAGCTCTTTTTATCTAGAGCTGGAAGGTTGCTGTCTTGGAGTG
GGGAGAAAGGAGACAAGTATCTGATGGGTGGGAATGGAAGGTGTGTATCCTTGCAGCAGACCTCCAGAGTAGCTGACT
GACTGATATGCATGGTAGTCCTAAGATGTGTTTGAGAAAAGAAAATATTAGGAGCTCTTGACAAATCTTGAAAATCAAT
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TGAGAGGCAGAGGTTGCAGCGAGATCGCACCACTGCACTCCAGTCTGGGCAACAGAGTGAGACTCTGTCTCAAAAAAAA
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GGTAGAGACAGGCAGAAGAGATTAGTTTGTATGTGTTATCCTGAAATCTGAGTTATTTGACATTTTTTAAGGCAAGGCAT
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TCCATATTATAAGAAGTCTTTTCTTCATATGTACCAATTCCAAATAAGTGGATGCTGGAAGAAAATCTGTAATTTATT
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TAAAATTGACAAAGGAGAAGTTCATATAATCTAATTCAAATCAAATGAGTTTTTCAAAGAAATAAGTTTCATCATGT
AATCTAAGCCAAAATGAGCTGAAAATTTACACACAATTTCTTTTCTTTTCTTTTCTCTATTTCCCATTAGGTGGTTTT
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GT
AGCCAACAGCAAATTAGAATTGAACTCCATTATGTCTCTCAGGTCTTCTGATCAGCTCTTATTATACTTACTCTTAGT
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TATTACTGAATCTCATAACTCACTTTTTCTCTCCATTTTATATTTTCAACTTTTCTAGTCTAGGGCTCTGCACAAGTTAT
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GAAGTTATTAACAAACGTTCTTGAATTATGTGTGACAGAATTTGCTGGGAGCTTGAATCAGTATATTTGTTTAGGCTG
CATATAACAGAAGGCTTCATTTAAATAGCTTAAATAATAAGAGATTTATTGAATTATATAATAAGAAGGTAGTGATAG
AATGATTTCTAGTAGTACAAACATGCATCAAGAATCCAGATTCTTTCTTCTTTCTAGTCACCCATCCTCAGCACAGTAG

Fig. 9.323

TTTTTGATCTCAGTTTTATCCTCTTATGGTCTCAAGATGACTACAGCAGCTCTAAGTATATTTATACAACCTTCCTTTTC
TTCTGTCACTTTTTTAAGAAAGATAAAAACTTTACCAGAAGTTCCTTAACAGATTTCTTTCTCAGATCTGTATTAGAAAC
AAATTCATCATATGCTTATATCTAAGCCATTTCATTGGCAAGAGAAATGCTATGAAATTGGCTGGGAAGTAATCATGGTT
CATCCTGGTCCTGGGAGGGGCTGATCTCCCTTGAAGCACCAACCGCCTGACACCTGAACAAATTCTAGTGGCTGCTGG
GTAGGGAACAACAGTGTCTTTACAAGTTGGTGCATTATAAATACATTTCCATAATTTGAATCAAGCCTACCATCTCCTC
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TCCTTAAATACACATTTTTGTACCTGCCATATTTTGCATGCCCTCTTTTATCAGCCTGGAATGCACCTTTGTCACCTTTG
CAGCCTGGTAAACCATCATTTGTGCTCATAATTTCAAGAACCACCTCAAATGTCTGCTCTTTAGATCTGAACCCTTCCTG
AATTCCCCAGCTTAGCAGGTCAAGTGAGTTGTTCTCAGCTTTGTACTTCTGCTATAGATCAGCCCTCATCATTTTATAA
CTATGTATTGACCCATTTACTCCTGAGAATCCTGTAAGGATGGGACTCAGTTCCTTGGTATTTTGTCTCAGAGAGATG
ATGCTTCATACTTACCAACTGTTCAATAAAGTTTGTGTAATTAATATAGTTAAAAGCAGATACCTTTCTGTTATAAGTT
CAAAGCCTAGTGATTGAAGATTTTTCTTTACATTATTTAAATGTATTTATTTTGGTAAAATATTTAGAGCAAGTTTCAT
TCCATCTGTTCCATCTAAGCGATTTCATGACCTGCATACATTCTAAATGACAGGAACCACAAGCAACTCAGTCAGCAGGC
TTCAGAATCGGTCATCCTAAAAGATATTGCAGGTCAATGGCTAATACACTAACAAAGAAAGAGCTTATGCAAATAAGTA
AGGAAAATACTGTTTCAATGTTAAATCATTAAAGACAAATTCAGACTACTCACAAAGAAGATCTATAAATGCCAAGTTA
GAATAGTAAAACAGAAAACAAAAAACAACATGAAGACCTCTGATTACATCTATAGTAATTAACATATCCATTTAACTC
CTCTGCCCTTCAGAAAACCATCAATGGATGTTTATAGGYATAAATGACAAGGACAAAGAAAGTGGAAGGGAGATAACAG
CAACAAAATTTTGAAGCTGAAAAACMAGTATACAAGGGGTATACTTCCTGATGAATGTAAAACCGTCCCTAGACAAGG
GGGGCCTARGCAGGTGAAGACTGAGTGCCATTCTGAAAACAGAGGGATTAAATAAACACTTATATATTTGTGTTTCATG
ATAACACAAGGCTTTTTAAATTCTGTTTACCACTGTGTCTCCGGTGGCTAGAATAGAGGCTGCCAAATATAGATGCTCA
AATATTTGTTGAATGAATTCGAAGRTTGAAAGTCCCAACCTTCTTCCCCCTAGTCAGCTTCCAACCTATGCTTAAGCCTT
CTGGACAGGAGATTGATATAACCTTTGGAGAATTCACCAACCCCAAGAGAAAAAACCCAAAGATAATGGTGCAAGGTATT
TCTGAATGAACTGTTTCAGTTGGATCATCAAAAAGTGGATGTAGTTGACAAATCTTACCTATGCAAACAGAGCTTCCAA
TCTGATTTTTTAGTACTCTAGCCTTAAATATGAGTAGACAGATAAGATTTATCAAATACTTAGAAAAAGTAAGTGAAAGA
AACCAAAACAAATGGAAAAAGACCAACTTCAAAGTACAGATGTTGGCCGGGCGCGGTGGCTCACGCCTGTAATCCCAAC
ACTTCGGGAGGCCAAGGCGGGGGGATTACCTTAGGTCAGGAGTTTCGAGACCAGCCTGGCCAATATGGTGAAACCCCGTC
TCTACTAAAAATATAAAAAAATTAGCTGGGCGTGGTAGTAGGCAACTGTAATCCCAGCTACTCAGGAGGCTGAGGCAGG
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AGACTCCATCTCAAAAAAAG
CAGAGAGAGGAGAAGAAATACACTTGATATTGTATTTCATAAATATTAACATTGCTTCCACGAAACAAAAGCAGATGCTA
TGGAAAAAAG
TCAATAGAAGGGTTGAAAGGTAAATTGAGGACATCTCTATGGAAATGGAGCAAAAATACAAAAGGTAGAAAATACAGA
AGAAAAAGTGGGAAATCAGAGGACCAGTTCAGGAAATCTAACATCTAAAATCAGAAATTCAGAAAAAGAGAAAACTG
CAGGAGGACAAAATTGAAGAAAAATTTCCAGAACTCATTAATTGGAGCATTTAGCACAAATATAAAGTATTAAAAAGAG
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TCGTTTTGAAACCTAGAAAAGAAATGAAGTGATGCCTTTTAAATTCCGAAGGAAAGTGATTCTCAACCAAGAATTCTATA
CTCAACCAACTATTGATCAAGAGTGAGCATGGAATAAAGATTATTTCTGAGGTGCGAGCCTTTAAAAAATGTATCTCT
GATGCATACTGACTCAGGAAGCTACTGGAAAACGTGCTCACCAAAACAAGGGAATAAACTCTAAAAACATTACCCTTGG
GATAATAAGGAAAGAACATCCCAAAGTGACAGCTGAGCAAGAGACATGGAGAATAACCAATCCAGGTCAAAGAGGCCTC
TGGATGAGATTTCTTCAAGAAGATGAATTTAATAAAATTTCTTGATGTGTTTGAGCCATACTTAGATTTTTGTAAATATGG
GAAAAGTTTGGGATTGAATTAGTGATAAGTATATATGGACATCTAAGGGAACAAAGAACTAACAAAAGACAAGAATTT
TCAAGAAGGAAACAAAGAAAAAAGGTAATCAGGGTATGTTACATAGTTTAGCTGCTTATAGTTTTTCTTTGGTTCTG
CTCATGGAAACACAATGACTATCAATCTAAGTAAGACTATAATATATTAGAAGGATGGGTGATGAGAAGTGTGAAGTGT
TGCAAAGGTAAATCCTTATCTTCCGCTATGAAGTATCAATAAGCAATGCCCAAAAAAATGAACCTATTAAGAAGTAACTG
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GAGTGGAAAGATACATGTATGTATTGTGGGTGGGGGATGCACTGCAGGAGATCTCTTTTTTTAATCCTTGTGGTACTACT
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AATACACATTAAGAGGTATAAAAAATGTTTATTTCTTCTGACTTAGTACTTCTGCTTCAGAAATCTCTTACAGTGATCTA
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AAGAAAAGTTATTTGTTGCAACCACAGTAGACCAGGTTAATGGTGCCAAGAGTGGAATGCGGATAAAGGCTGACAAGGC
CATCTGGAACGTGTGAGTCATTTCAGAGCATCACAGAAGAGAGATTTTCTGCAAGTACTAGCTGTGTTGACTGTGACTGCT
GTTTTCTGCTGCTCAGGAATCACCAGGGAAGGAGAGGGCTTCCCCACGGGATTAGGGAAAAAAGCTTC
TTGTTCCATAAGAGAGTCTACAGATAGGATGAGTAATAAGGGATAGATTTTACTAAGGTAGAACAAAATGTTAGGACGC
TGGTACGAGCACCCTGAAATATCCCTATATCAAGTTTTAGTCTTTTCATTGCATCTTCTGAACCTGCTGGAGATGCTT
TCACATGAACGTATTTGCTATAAACTTTTCTTTATCTTTTGTTCATGCTGTGAAGTTTGCTAATCTTAATGAACCAA
GTCTCTTCATGCTGACAACCTCATTGTAAAAGAGGTAAGTGTGTTTCCATGGTATGGGGAATGGAGAGGTATAAGGAG
GAAGATGGATTTAAATTGATTTTTTGAATGCTTGCTTTATTTTATCAGTTAAAGAAAAGGTCTAACGGATTATTTAGAT

Fig. 9.324

AACTTTAGGCTCCAAGTGCAGTCTGTTTGTTCATTCCTTTAAGAAACCATTTTGGATTCTGACTACTTGAATTTGCAC
ATTTTGTTCCTTCTCTCTGGAATACCTTAACAAATTCCTTCTGCCTGATAAACTACTTACTCTTCCAGGTCAAGTTGAA
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GTTTACCTAATTCTCACACCATTAGATGAGCGAGGTAGTTTTAGGCTAACTTTAAAAAGGAGGAACTGAGACTTACAA
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TGCTCATCACTGAATCCAGCATCTGTGAGTGTCTGCCATATAGGAATTGCTCAATATACATTTTTTGGAGTAAATAACTGA
ACTAACAATGAGTGAACAAAGAACAATGAATAAAAAACCTCAGATCTAGCCCCTAACAAAAATACTTAAAAAAA
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TTAAATCAAATGAAATCTATATCCTCTGGGAATGCTTGTGGTATGGCTTAGGGACAAGCTTTACTTATGAACAATGATA
CTGAGACTTCACAATAGTCAGCTGTGAGATGTGAGACTTTGCATTTTACACATGCTTTTAACTAGAGCTCAAATAGG
CAGTTTTAAGCCCTGGACCTCAAGTCAATGTGGTTCATGTTTTGTCACTTCAAGATCTACAATTGAACCTTCATTACGAT
AGTCTTAGATGGTTTTTTCATAAATTTTGTAGTCATGAAAACTGACAACATATGAGTCTCCAAGTACCTTTTAAATATATG
CAATATTTTACCTACTTAATKAATACATGTGTTTTATTTGATAACTAAAAAGTTTATAAAGTCTAGAAATAAAGAAAAGT
CCATGTCCTTTTTTCTTTTTTGTCTTTTAAATGAAAACCTCATGAGAAATAAGAGGGCAGAATGCATTAAATTATTTTC
TTCTGTAAACAGCACAAATCTATATCAGATTTTAAATACAAAAGAACATGCAAAGGGATAACAGCATTGACTTCAGTTCT
AATATAAATAAGGCAGGAAATTGGGTGAAATCAGTTTTTCTCTAATCTTACATGGAAAAAAATTTGTCAATTTGGCAAA
CCCATATGAATCCGATCTGTTTGGTTTATTCATCCATGCAGTGACATTCAGACTCCAAAAACTGTATCAGAAACCTAGT
CAATATTTAAATGCCATCATATAGATTAGAAATGGAATAAGGTATAGGTAACCTTACTGCATTTCAAAAAAAGTACTAAT
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GAAAGGCCTATTTTTCTTCTGTCTTAGTCCATTCAAGCTGCTGTAAACAAATATCATAGCCAGGCACAGTGGCTCATG
CCTGTAATCCAGCACTTTGGGAGGCCGAGGTGGGCAGATCACTTGAAGTCAGGAGTTTCAGACCAGCCTGGGCAACAT
GATGAAACCTAATCTCTACTAAAAATACAAAAAATCAGCCAAATGTGGTCACAAGCACCTGTAATCCAGCAACTCAG
GAGGCTAAGGCACCTAGAATCACTTGCACCTGGGAAGGAGAGGGTGCAGTGATCTGAGATCATGGCACTGAAATCCAGCC
CGAGTGACAGAGCAAACTCTCTCTCTAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATAAATCAGG
AGCTGGGTGGCTTATAAACACATAAATTTATTTCTCACAGTTCTGGAGACTAGAATGTCCAAGGTCAAGGCACGGTAG
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GACTATAGTATCTGCTTTTTTGCCACAGAATGAAGAATAAACCACGTCTTATAAATAAAAAATCTTTCCATGGTCCCCCAC
CACATAAGGACATGACCAACTTCCTTTGCTAGTCATCCATCTCTCATAATCCTCTCCAGCCCATCTCTGCAGCCTCATC
TCCACCCACACATGCATTCTGTGTAAACAAATGGTGGGCAGGCTTCCAAATGTACTGTGTTCCTCATGAAGTTGCACCTT
TGCTGATGTTATTCTCTTTGCTTGGGTCCCCTCCCCTATCTATTCCTCTCTTCTGCACTTCACCTTGCTCTTCTTGTG
CCCTAATAGCTCCTCCTTTTGTGACACATCAAAAATTGCCCACTGTAGGAAGCCCCTCCAAGACTAAGAGTGCCTCTCT
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ATTTGTTCAATGAATCAATGAATTGATAGGTTTTCTTTTAGTAGAAGGAATAAGGTACATTTTTTTTTTAGAACAGCATCC
GAAAAGCAAGCAGGAAAAATTGGGACATCTTTGCTTAAAGGTAAAAATGCTTTTATGGGGACCACCTTTGAAACTCCATC
TACTTGGATCCTTTTAGTCTTCTTTATGAGATAGGAATATTAATTCTTAGATCCAATTAAAGAAGGTCTAGTTACCAAG
GAACAAAACAAATTGGTATAGAATGGACTTTCTTATAAGAGAGTCCCAAATCATAGATCATAAGGACAATCTTATGTTG
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AAGTTTTTCAATTGTACTCCTTGACTATTTTTATTGGTGTACAGTCTCTTGACAATTATTTTTCTCATCTTCTTTCCCAAGGA
AACTTTAGATTTAATATGTTTATTCTGCAAGATGTTTGGAGACAGGTATACACCTTAAACACCATTACAATCAATGC
CATAAACTTATTCATTGCCTCCAAAAGTTTCTTCCCATCTTTTTTGGGTTTTTTTTGTTTTTCTTTTCTATGGTAAGG
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CAGCAGCTTTCCAGAACTTATTTTGCAAGACTAAACCTTTGTAACCTTGACCACATCTACCAATTCCCCTCTTCCCCC
AGCCTCTGGCAACTACCATCTACTTTCTGCTTCCATCAGTTTGACTATTTTGGATTATACATGTAAGTGAGATCATGC
AGTATTTGTCTTCTGTGTCTGTTTTATTCACCTTAGCATGATGTCTCCAGTTCATCCAACCTTGTGTGTAATGACAG
GATTTCTTCTCTTTTTTAGGGCTGAATAATATTTTATTGTATATGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTG
TG
TGTATCTTAGCTATTACAAATATGCTGCAGTGAAAATATCTCTTTAAGATCCAGATTTTCAAGTTCTTTTGGATATATACC
AAGAAGTGGGACTGCTGGATCAAGTTACTTCTGTTCGAAAGTAGGGCACCCTCTGGACATTTCTACAACCTGAAGTGA
TTGATTCTTACTTTTGAAGTCAGGCTATTTGTTGGAATGGTGGGGAAAAAAGCTAGAGCTTAAAAATAAATCTAAAGT
TGGAGAGGGGATTATTACTGATGGATGCATTAATTCAAGGGTACTGATACTCTAAATACATTTTTTAAAGTTGTTTGA
TTAACAAGTCAAAACAGATGGTTTAATGACATTTTAGAGAGGTTTAATAGAGATCCAACCTGAATTAACAAATCACCATG
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GATAAATCACTGGGAAACGGGTAGCCTGTAGGACATGACAGCAAACCACACTTTGGCAGGACCAGCATCAGGGCTGCGT
GCTTTTTTTAGGAGGCTATGTAGTCTAGCTAGCACAGAGTGACCCAGTGAGAGCTTGGTTGGCTAGGACTTTCTGGTGGT
AGCCACAGAGCTCACCTGAAAGAGTACCAAGGATGAAAATATCATCTTGGCTAATTGGTCTGCTAGTTGATTTAAAAA

Fig. 9.325

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ATAAACAATAAAAAAATTTCCAGTGTATTTTAGCAAAGTTTAATATTTTGAAGGGGGCAGAAATGTAGCATATTTTGG
GATCTTAAATCTGAATTTGTGTCTTTTCCAGACTTTGTGTCAGTGTCTTTGTCTCTCTCCCTCTGCCCCCTCCATGTC
TCCCTCCTTCCCTCCCTCCTTCTCTCTTTCTCTTCCCTGGTAAGTATACTAGAAAAAGTATATTTTTTCTCATTTTTT
TTAACTTAATAACAACAGCAACAATGACAACAGTCAAACCTCTGAATTCTGGTCACAATCCAGATCAATAATTTTTTCC
TTGTAGTTACCTTGGAATTAGGTTCTTAGCCCCCTCACTGCTCTCTGTATATTTCTGTACAATATTCATCAGTTTAAATA
AGTACTTCAATTTTCATACCAAAGTGCCAAGACTATCCTATTTTCACTCATCTTTAACTTTCTTCTTTTTTCATTTCCAC
ACCCCTGCCATTGCCAAACAAAAGTATTATCTCTGTTTTTATCTTTCTTTTAGACTTTTTTAAGGCAATAAAAAATTGG
CTTGCTCAGAAGGCTGATACACTGAGTGTTAGAATAACCCAAAGGTTAGAATGCTTTGGTCAACTTATGGGTATTTTTG
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TTACTAGCATGATTACTGATTAACTTAGACTGTTGGCTTGAAGTTTAGTAGCCTGAAGGAAAATTTCCAGAGGCATT
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ATTCAATTATGCTGCAAATGCAAGTGATTGTAGTACATTGGCCAATAAAAGTGAAATCTGTCTCAACAAGAATGTTGA
AAGAAAGAAGGCAGTGAACTTTTCATCTCCACTGGGAGTGAACCTGCCAAATCTGGGTCAATCATAGTAAAA
TGTTCAAAACAAATAATGCATCTTTTCATGGTTCCTGTATTTCATCAGCACTTTTGAAGTCACCAGTCTAGAGTAGTCGC
CTTTGGCCAAGTTTTTCAGTAGCCTGTGGTTTAGGCAATAAGCCTCAACTGTCTTTCTCGAGGATATGTTCCAGGTGGT
TGTAATCCATGGGCCATTTTCAATCAAATCTGAAAGGACAAAAGGGCAGTTCTGTTTATATGAAATGACATCATATTAT
AACCTCAAAATTGTTGTCTTAAAGTCAAATCATTATCTAAAAGGTCTCTTAGAATTACTTAGAGCTTGAATGCAAAAG
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AAAAAATAAATAAACCACATGCATTCACTCATTTGTAAACATTTCCACCTCATTTGCAGTATCACTCTATGTGCACATAC
ACACAAATATACATACACACACATTGTCTTTTTTTCTCTGAACCACCTGAGAGAAAAGTGCACACATCATAGCCCTCTA
TCCCTAAACTCCTCGATGTATTCTCCCAAGAACAAGGACACTTTTCTGTATAACTATAGTCTCAAACATCTGATTGCC
TTTGAATTTTATAGCATGACTTATTTATACCTTTTTTACTTCGAGTTCTAACTCAGATACTGAGCTATAGAGGAACAGCT
CAACTATTAGTTGAGGGATAGCTCAGATGTCTGATGGAAAAAGTCACATCAGTAACATTTTGAATTTCTGCAAATATAC
ACAATATGCCAATGTATGGGAGATAACTAAATACTGGTTTTAGAGAAAATAGATGAGTTAATACAAATAGAGATTCCCC
TTAATTAATAGATCTTATATGTTTGGTTATATGTCTTATCCATGCTATTATAGCTAGCTAGCATTTCATTAGGACAGCCA
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GTTCTTACTTAGATAATGCTATTGCCCTTGCTCCTTAACAGTAAGCAGTGGAGAAGCAGAGTGGACGCTGGTGATGCTGC
CTTTAGTCTTGAGGGAGTGTCACCAGTGGGTAATCATTCTGACCCTGCAACACAACCATCCCTTCACTATTTTCAGTG
AAGGTCTCTACTTCGGTGGTAGGTCTCTACTCAACATGGTACTTATAGCCGTTTTTATACTTGCTATCTTAAAAAATAT
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TTGCAATTTTTTTTTCCCAAACGAGTTTGAGCATGCCATCCCTCTCTACCCAAGTATATTTAACAGCATTTCTATCTA
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GCTGGAGCAATTTATGAAAATTGGGTCCCTGGTAACGTAGTGSATGTTATACAGACAGTCTGTCAGAGTCATCCACG
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TCTTCGTTTTATTCAATTTGACAGATTTATATTGAGGGGCACTGTTCTAGGAGCTGGGGATTGTTGTAATAAATAAGATAGT
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GACCATATGACTGCAATATGTCAATTTGATAAACCTTTTCAGGCATGACTTGGACCCACCCAGAACTTAGCTTTAAAGCTA
TCAGATTGACAGGAGGAGAGAACACTGGCTTTAGATGACAAATCATGGAAATAATTATGAAAAGACCATCCAGTTTTTA
ATGACTTTACAAGAATAGTGTTCTTGTGAGACTATTGAATAAGAAGAAATATATGAACATTTGTATCATTCAACTGTCT
AACTGATACCAACACCAATAAATATATGAACATCCTTATTATTCAAGTGTCTAATGAGTGAGGGACAGTGCCTGTAGGA
AGGAGAGCCTCCCAACAATGGAAGCAGTGTTATTCTTCATGGTAGGGAGACCATTTTTTTGGTTCCTCACAATGACTTCC
AGGCTGGAATGATTTCCAAGTAGGGCAGGGAAGACAAATATCTGAAGCCTAAGGGAAGCTCCAGGGTGATTCTTGTAGT
TTTGTGTTGGCAGACTTTCTAAAAAGCATTGTGAGATAGTTTTGGAGCAAGCAGTTTAGCCATCTACTACCACACAATAT
ATTCAAATCAGAGTAAGCTGGAAGAAAGTCATCCAACATCCAAGTCTCAGCCTGGCCTATGTGAACATACTTCTTCTTA
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TTATTGAACATCAAAGCAAAAATGGTGAATGAACAGATATATGACACAAAAAAGACTAAGGTATAATCCTATATCATGA
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AGTGGCTTAAGGAAGACAGATATTTCTCCTTCATGTAAATCTGAAAGTTGGCAGTCCGAGGTTAATATGGCACCTCTA
TCCAGTACAGTCTTCAAGGACACGTGCTCTTTCCATCTTATTGCACTGTGAGGTGTAGCCTCTGTTGCTAAATTCAACT
CACCATCCAATATGGCTGCTCCGACTCCAGCCATCAACTTCACATTCTAGCCAGTAGGAAAGAGAAAACAATAAATACA
GGCAGATAAAAAGATAGATGACAAATTTCTCTTAAGGGAAGTTTCTAGCACATCATATAATATCTACATAACGCTTCTA
CTACCAACTCATTACCAAAAAGTTAGTTATATGGCAACACCTGGCAACCAAGGAGGCTGTGAATTTTCTCTATTTTGGG
CAGTCACATGCTTACATGAAAACAGGTATTTGAAAACAAGTGATGAAGAAAAGCATCAATATTAGGGGACATTTAGCAG
TCTCTGCCACAATGTTGCCTATTAATAATCCTGCATACATTTTAAATATTTAATATCAGTCTGCAACACTCTATTTGCA
AGGTAATGTATAGTATAATCTTTACCATATGAACCTTAGTAGCCATGGTGTTCAGGAAAGTTGTGTTATTTTGTGCTAGA
ATATTTTACCTGCCCTAGGTAAGGGCCTAAAGATAAAATGTGCTAGACTAATTAATTTTAAATGGCATATAGCAAGAGAT
TTCCAGCGTTAAAGATTGCATCTCCACCTGCAATTTGGGAAAGGAAAACTGATAGCACAAAATAAAAGTAGTGGGTGT
CCTGAAAAGTGTCTGATGCTCTTCATGTTCTTACCAACTCATATCCTCATCTAAATTTGAATCACAGGTACATTCTGAC

Fig. 9.326

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CTTGTCGTAGAGAGAGTCTTAAAGTCTCACGAGAGAATTACCGTGCTTGTCTAACAAGACTGTGTTCCAGCAGGCATG
CCAAAATTTGATAGCTACAAATAGGGAGATTTGAAAGGAAAGAGGTGGGAGAGGTAATTTCTTTTAACTCAACTGCATC
TGGTGGATTAAAGGTAGACAATATTACCATTTTGTCTGACAGAAGGCAGACATGGCACTAAAGAGGGGAGAATGAGCAACCA
CAGAGCTGATTTAATTTCCAGTGGGGTTTATGGAAGCACCAAAACATGATGTAGCCAAATGTTCTTAAAGTATGAAGT
AATTTAATTTGTTCCATCATTACAAGAAATTAAAGCCAAGCACAATTACATCCCAGTAGTAAAAGGAACCCGCTGAATT
GCCATTGTTATTGACTAAATGAAGTGAGATTTCTGATTACCTTGTCCAGCACATTTTAATTTCTGTCTGTCCATTTATT
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TTGTTTCTGATGTGCAAAAGAGACCCCAAGTTTAAAAGGAAGACTTAACTTACGGAAGTGATTTTTTTTTTTCTTCCAC
CAAGAGTCTCTTTGTAGACAGGTGTCTGTTCTGTTGGGAGTGGACTTACACCTCCTGAATGCTGTGATTGAGAGAGCTG
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CAGTGAACCCCTCTCATAAGTAGGGTGTCTAGATGATTTAGCATTCAAACCAGAACACTTTTTAGAAATGAAAGGCAATGC
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TCAACAAACATTTATTGGATACCTAGTATATGACAGCCAGTGTTTAGCACCAGAGATCAAAAAATGAATTCATTATGGT
TCCAGCCCCAGAGAAATTCAGTCTAGTAATAAACACATAATTGTGATAGACTGTTTAGTGATTTAATAACTTAAAGAGT
TAACTTCTGATTTGGTTCTCATGCATCAAACATAATATTTGCCAGTCTCTATCTCTACAAGGAGCCCTGGATTTTTCCC
AGTCCCCTACTAATGCTAGATAATATGGCAAAATACACAGGCTGATCAGGCTGTTTTAGAGACTCTTTTAAAGCAGAGAT
CTTTTGTTTTCCAGACTGCTAATTTATTTTTTTCTACCCAGAAAGCCCTTCTTACCATCTGAGCTATTCTGACCAAATC
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GCTTTCCAGGATTACTGTTCTTAACCACATTGATAAATGCTGGGAAGACTATCTCAGTTATCCAGCATTGGATAACAGA
CTGTTGGAGAAGAGTGAAGCTTCAGTGTGAGCTGGGAAGATCCCAAAAATCCTAACATGCTCTAGGTGCCTGCATATA
AAATTCATCATCATAGTGACTTTAGGACCACTTGCTATTTTTTCAGGCACTGCTTTACATACATAGTCCTATTTAAATAA
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ATAAGGTGCCAAGCTAGTCAATCAGTGGCAAAGCTCAGATTTGGAACAAGGACTGCCTTACTCCAAAACCTGTTCTCT
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CAGGTTTCCGTGAAGGAGAATTATGCCAGCAATGGTTTCTCACCTTAATGAATTCATTTCTAACCATTTCTTGCCCTGC
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CTGCACATGTATCCCAGAACTTAAAGTAAAAAAAACACAAGAAAAACAAACAAAAAACTCCTTTTTAAACATTG
TCTCAGCGTCCAGTCCGTTGAGTTTCTGCACTTTTAGACACTCAAGCCCCCTTCTTTCGTCTTTGCTCCAGTGGTTGTA
GCTTCTTACCTCCTTCTTACCCTCACCCCTAGGCTTCCACTTTAGAGAATTTGGAGAACTGGGGAAAGAAAAGACTGTT
TTGTGCATGGTGTGCCTGCAGCCTAGTGCCATGAGCTGGTTCTTGCCGTGTCTGAGCCACCAGATCCACTCCTTCTGTG
GTTACACGCTATTCCCTCTATCCCAGCCAAGCTTGCAAACTCTGATCATGAACCTACCAGGTGGAGGSGGGGCATGGAT
ATCCTCCCCAGGAAGCCTGTCTGGAACCTCTGAACCTTTCTGACCTGTGCATGCCTCTACCACAGCACACCTTAATTC
TTTTGAAATCACTGCATTTGTTTCAATTTGTTAGTTGTTTGCTTCTAGATGGTAGACGCCTTGAGGCCAGAATTTCTTCTGA
ACTAAGCTTTGGAAGTTGAGCACCTCATAGAATACCAAGCTTCTCGTATTACTGGTGGGGTTCAACAAGTATTTGTTGA
GAAGTGATGAATCAAGCTAAAGTTTAAGTAGGCAAAGATGAATCACGTCATAACCTCTGGTTTCCYAACTTTGTCTATA
GAAAAGGCTCCTTGCTAGGTAAAACAAAAACAGAAGAAGTATGTGGATTCTCTTCACTTTGGGACTCATCACCCCTGAA
TCTCTCAGTTATCTCAAGACTGATTCCTCCACCCTGGAGGGCCTCCCTGCCCTCCTTTGTACAGCAGTGAAAGGCAG
AAATTGAATAGAAGGAGGGAAGAGGGGAAAAAGGTCTAGATGATCAAGTGGCAAAAACAAATAACCCAGCATGAACAAG
TATGCAGAGAGGGAACTCTGAGGAACTTGCTGACAAAAGATAGAGATGGAGGTGAGGTCAACCACAGGGGAATAATGG
GGGCAGACAAGTCTAAGGAGGTAGATTTTTATAGGGACTTCGAATAGATGAACCTGAAGTTTGGGGAAGACCTAAAGGCAT
TAAAAATCCAGTATAAGTTCTTGATTAAAGGACAGATATCATAATAATTATTATTATAATAACAAGGGGTGCTTTGGGAAC
ATAGGCTATGTAGGAGGGATTGACAGAGGAAGGGAGGAAAGATCTAGGAAAATGAGGGCCAAACGGAAAGGTGTCTTC
TCCTGTTTCAATTTGCCAATGAAATGCCTAGGATATATGGTGGTCTGCTCCTACCCCTCCACAGTTCTAGCCACTGCAACCA
GGATGCCCAGCAACCATGTCAATTCATTTCATATGGTCTATCTGTCTCCTATAGGCATTTTCAGTGAGTGGCCGCTGC
TTTATGTGTTCTGTTGGTAAACTTGTAAGTACCCTGAGAAAACCTCACACATTATGGAGAAATTACTTCAAAAAATATGCA
CAGTAGTTAACTGAAATCTTTTTATGTGTTCTACTCTCACGTGAAGTAGAGAAGTAGAGGGAGAGTTTTTTAATTATAA
AAAGGGAGGAAGAGGGGAAGAGGGGAAACAAAACCTAACATTTATTAAGCAAGGTAATTTCTTACCTCAGTGTTTTCAAACC
AATTGGTAGATTTCAAATCAATTTAAGGGGTCACAATTAATACTTTAAATTAATAWAAATAGAAAACATCAGAGTGA
ATCATACATAGTAAGGATAAATGTTATTTTCATGTTAAATTATGTTTCAATTCTCTGTATTGGCTCGAATATATATTTAC

Fig. 9.327

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TCTTATGTGTATGGTATATACTGTTCCCTATATATACATTCCTTTTTATTTTATATATATAAATTCCTTTTTATTGGTTTA
ATTTATACTGAAATGCATGGTTTGTAAAGACTGTAAACAAATTTTGCCATGACTATTATTTATAAAACACAAGAGTTT
CTTTAAATAGGAGTCCACTGCTCTCCTCTACCTTTTAAAAATAAATATAGGTATAAAACCTTTTTGTACATTTTTTTTA
TATTACTGAAAGCCAGCATCCACATGCCTTCAAAGTGAGCACCTCTTACTTTAAACAAATCTACCAGGTTATAGTATA
AAATATAAATTTTACTGTAGTTTACTGTCAATAAACTTTGTAAAGCCACTGCTTCATTTCAATTTAAACTTCCCCAAAACC
TTGTATGAGTGGAGTCTGGGGATTTCAGAAGAAGACGGTGGTCATCACAGAAGTTGAAATTCTCTACTGTTTACTTCTTT
TCAAAGGTCTTGTTAGCGGTGGCCTTGAAGGTGATGAAGAAAGACACTTTCAAGGTGGACAGGTATAATTATTGGGTCA
CCAAGAACAATTTGAAAAACAAAATATGCCAATTGAAATTAGCATGTAGCACCTAAATACCCAGAAGCTTCTTCATTG
AATAATTTATTTCATTGAATAAATTATGTAAACTGAATTAGTAATTCCTAGTGAAATAGTGGATGAATTGAGAATAGTGG
TATTGGTAAAAATGGAAAAAATTTTTTTCTTTTGAGACAGAATCTCACTCTGTCCCTGAGGCTGGAGTGCAGTGGCAC
GATCTGAGCTCACTGCAACCTCTGCCTCCCAGGTTCAAGTGAGTCTTGTGCCTCGCTTGAGGCACAAGATTAGATTTAG
TAGAGATGGGGTTTTGCTTTGATGGCCAGGCTGGTCTCAAATTCCTGGCCTCAAGTCATCTGCCCACCTCGACTTCCCA
AAGTGCTGGGATTACAGGCATGAGCGACCGTGCCTGGCCAAAAAACAACAACAAAAAACAACAAAAAACAACAA
CAACTTAATTCATAAACTTAATTCACAATTTAAGAGTTATACTTCAAGAATGAAGATATATAATGAAAATATGAAGT
CTTCAGAAGGACACACATATAATAATTTTAGTAAATCTCAAATTTGTCTCAAATCAAAAACCTTGCTACACATATTTTTG
TCTTAATTATCATAATTAAAAATGTCTACACATTACATAATACTATCTAGTAAATATGCCCTAGAGATTGTGTGTGTGT
GT
GCAATTGTTTTTCTTTTATCAAGGAGCTGTCAATTTAAGTGCAAAGCTGGAGTATTCAAATAAAATTTGGGCTTACAAG
ATCAGGGCATGGCAAGACCTATCCAGATTGCTGTGGAATAATATATGCTTCCAAACGCTAATCTTAAGGATGAAAAACA
CATAAATGACATGCCTTCTTCCCCTTGCTTACCTTAGTGACTACAGTTTAAATAATCTAAAATAGCAGATACCATCAAT
ACTGTTGCTAATTATGCTACTTTTCAGAGAGGTACAAGAGTCCTTGTACCTCTAGTAGTAGTGATACCACTACTAGTAT
CAAAAATACAAGATTGTTTTGTAAATTTTGACCATGTTTTAAAAAAGATAGGGAGGAGATGAATTACTATTAAAAGCA
GCTCTTATATTTTACATGTAATAGATGATAATTACAAGTTTTTAACAAATAACACATTTAAGTTGAACTGAACCGCCAA
AAAAGATAGAGATATGAATATATGATTTTTTAGATTAAAGTTAGTTTTAACATTCAAATAAAGTGTGGGAAAATGGCCTG
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AACTGTCATCGTAGACCACTAAAGTGGATACCTCCACAACCTCTTGCTTCCACTGCATCTGGACTACCACAATAGCTC
CTTAGTCTCCATCCTTCTCTCTTCTCCATTCTATCCTCTTAGGCTACCAGTGCATTATCCAATTCCCACAATATCCAC
TAGCTCAAGGCCACACATTTAAGTGTCAACTCTATATCCCTTCTCCTCCCTGGAATGTGTCTACTCCACCCAGCCCAT
TTACTGTCCACACATAACCTTTATTTCAGGCTGCAGACATTTCTTACACCTGAAAACAATGCCAGCGGACTCCCTCCT
CCCACAGCCATTAAAACCACAGTCTTCTTGAAGACCTACATCCAAGTTCACCTTCTTCTGAAACCCAGCACACTGAG
CTATAGTCCATACCTCCCTCTCCCTGCTCGAACTACAGTAGCACTCCCAGGCTGAATGATTTATATGCTACCTTACACT
GTCTCTAGGCAGCAGCCGATCATGTGTCTGATTTTCTCAGCAGGTTATTATGGTTCTTCTGAGGCTGCTGAA
TCTTTTCATGTTCTTGTATTTTACAGCAACTATCACATCAGAGAGGCTCAATAAATATGTATGGATTTTTTGGACTATTCT
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GTTGTTGTGTGAAAAATGCATTCTAAAATCAAACAGGTTTAGATAATGCTTGTAAATAAAATTAATATCTCATTTTGC
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CTTGTTTGACCACAAAACGTCATTAGTAAGGAGAATCTTATAAGGACAATATTGATTGGAACATTCTTTGGGAAATGTT
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GTTGACTTTTTCATGATTTGGTGTGGTCTTGTAAAAGCTATTCTATTTTGTGTGTTTGTGTTTTTGGAGACAGAGTCTCA
CTCTGTTGCCCAGGCTGGAGTGCAGTGGCAACCATCTCCTCCACCTCCAGGGTTCAATTGATTCTCCTGCCTCAGCCTC
CCAAGAAGCTGGAACCTACAGACATGTGCCACACACACCTGGCTACTTTTTGTATTTTTTAGTAGAGATGCGGTTTTGCCA
TGTTGTCTAGGCTGGTCTTGAACCTCCTGGCCTCAAGTGATCTGCTCGCATCAGCCTCCCAGAGTGCTGGGATTACAGGC
GTAAGCCACTGTGCCTGGTCTAAAAGCTATTCTAAATCATAGGTTGTTCACTCACATTTGGCTCAGAACTGGTTAATAT
TTGAGGGTTGATGGTGATGATGGAAAAATAGTAAGTGTTTTAGGATTGAGAATCTGTGGAAAAGGAAGGAACTCTCAA
ATATTAACCTACCTTATTTAAAATTAACCTATTTAAACAACCTCTGCAAAGTGGTTCAGTATACCTACTATTGATTGAATA
CCTTCTATATGTGATACAGCCCTTGGTACTTTTACATTTCTTATGCATTTCTCATAATAACCCTGTAATTCTCAATGT
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TGAGTGAATCAAAGTCCATGCTCTTTCTGATCTATCACACTATTTCCCATGAAGAGCTCTTATAGGTTGTGGATTCTT
CTGTGTGTATAATAACTTTCTTAGCCAAATCTAAATCTCCATAGATAATCTTGTAAAATTTATAAACTAATTTATCTTA
TTCGTGTATGGAGCCAGTTTCATATACAACTGGATAAGCCAAATATAACCGATATACTCTTGAGTTCTGAAATTTGTTCT
CTATAATGCACACACAGTTAACAATGTAGGTTTACCAGTAGGCCAAAATAGTTTATCACTCATATGTGTGCTTGTA
TGCAAGTAATGAGATTAAAAATGTACATAAGAAATTACCTTTCTGAGACTCTGTTTCATAGCCTGTTTAAAAGGGCCTA
GCTTTTTTCRGGAAATCTTTTGTGTGTTTTTTCTTTTCTTTTTTCTTTTTTTGGGACAGAGTTTCGCTCTTATTGCCCA
GGCTGGAGTGCAGTGGCACAACTCTCGGCTCACTGTAACTCCGCTCCCGGGTTCAAGTGATTCTCCTGCCTCAGCCTC
CCAAGTAGCTGGGATTACAGGCACCCGCCACCATGCGTGGCTAATGTTTTGTATTTTCTTTTAGTAGAGATGGGGTTCAT
CATGTTGGCCAGGCTGGTCTCAAACCTCTTGACCTCAGGTGATCCACCTGCCTTGGGTTCCTCAAAGGCTGGGATTACAG
ACATGAGCCACCATGCCCGGCTGTGTGTTTTTTCTTAATCCCAGTCTTCAACTGGACAAATGTCTCTTTGGCATTACTT
CTTAACCTTAGGTATCTCATAGTGAGAGAGATATAAATGCTAAACAGAGATTTATGAGAAAGTTAAAAAAGAAGAA

Fig. 9.328

GAAGAAAGCAGAGATAGCGATTCCGAATAAAAGATTCTGGGGCAGTGTTGATTAAAATAATTGCTTTCTTTTCTTCACT
CAGGAAAGTATTCTTAACCTTGGAGTCCTTGGTGACTTCAGGGAAGTCAGTGAACACTTTTAGAGTGAAAAATATTGATA
ATATGAACTTATGCTCATTTTTCTGGGGGTGTTTGCATCAGATGCACCTTTGTGCATTCTATCTGTTTCTCCCAAACATC
ATACTGGTGTATTGGTGAATTCTTTTAAAAATTTACTCTATTTTGTAAAGGATCATTACATATGAATGATATACCATAAT
ACTTGTCAATTTTTATTTTAATGTTAATATTTCACTTCAGTAAGACACCATGATCTGCTTGACCATTACCAAATTTTGGC
AATGTTAGTTCCTAATACTCTTTTAAAAAAGAGAAAGAGTTTGAAAGCAAAGACTGAGAACAAGAGATAGACGAGGG
TGATTACATGTAGGAAGCCACACCCAGGCCAGTATTACTGTTTGAATCTCRTTTGGAAATAAATATTCTTATCTGATAG
AAAACAAGCATACTTACTGATTATTCACTCACAAATATTTGCTGAGTGCCTGTAAATGTCAGGAATTTTCTAGACAGTT
ATAGAAAGGCCTAGACACAAATATAAAAATGACATTAGAAAAGTCATACAGGCAGAAGCCAGCAAATTTATTTCCAGGTG
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GAAAAAGGCCATGTGAACAGAAGTGTGACATTAATAACAATAAGTAGGAGAGAGTTGTGCAGCAGGTTCTTAAAGAAT
GAAACAATAAAACCAGGGTTTAAGGAAGATTATTCTGACTTTATAAATAGGACTGTGTTGAGAAAAAGTGAATCCAAGG
AGATCCAGTAGTAGACCATTATATGAATCTAGAAATACACAGATGAGAATTTGACTGAAGGTGACAGTTACAGAAATTA
AGAAGAAATTGGTAAGTCAACTGGCAAAGTATTGGATGGGAATATGGTAGGGGTAGGAGTGGAGATAAGCAGAGTGAGT
GTAAATATACTCCAGAACTTTCTATTTTATTAATAGCTTTATTTCTGAGAATACCTCAATTTCAAATAGAAAACATGTAC
CCCTGAAGAACAGTTGAGCTAAACTCACAGAATTCAGGATCATGGTATTGGATGGGATCCTTGACAAGTAACTGGTCA
GTTTAGGAATTCCTCTACAAACATAGCTGACCCCATCCAGGCTAACTATAATGAACAACCTTAAACCAACACCTAAC
CCAGATAGTACAATTTCAGAAATCAGATAATAAATGAATGTTAGAGCTGGGTAGGTGCTAAAGATAATCTGTTCAAAC
CCTTGTTTTCTCTAAGGAATACTGTATGCAGTAATTGACAAAGGTGAAGAACAAAAGACTGTTATATCCTAAGATTGAT
AGTATATGATGGAAACACAGATGCTCTCCTACAGTCCCCCAGGGAAATAAAATTGATTCCCTAAATTTACAACAAGAAT
ACGTAAATAGCATACTAATTGTAACTTTTTTTTAAACCTGTTTGCAAAAATTTGTCTGTAACCTTTGTTCAAACAACCTTC
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AAGCTGCTATAACAAAATACCATAAACTGATGGCTTAAACAACAAACWTTTATTTCTCACAGTTCTGGGAGCTGGAAGT
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ACATGGTAGAAAGAGAGCAACCTCTGGCCTCTTCTTATAATGGCACGAATCTAATTCATGAGGGCTCCATTCTCATGAC
CTAATTACTTCCGAAGGGCTTTCCCTCCAAATACCGTGACACCGGGGATTAGATTTTCAGCATATGAATATTGGAGAGAC
ACAAACATTCAGTCCATAACAATATTTCCCTTGACATTTCCTTTTCTTTGCTCCTGAGAGTTTTTCCTCCTAACTTT
CTGCAGCTTCTCCTTCCCCTTCAGGGACTGCTTAGGATTTTCTCCTTATCTGGTCACCAACCTCTTAACTGGTCTTTAC
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CACTCTTCCAGTTTCTTTCAAAGAATAATATCACTCACAGTTGCACACATGTCACATGAAGCCCCAACCTAGATGCCT
AATTAACTTAGCTCAAACCTCAATTTTTTTGGACAAAAGGCCTCCTTATTCTTATAAAAGCTTTCTCCTCTTTCTTTGA
CTCTTCTCTTATGTGCTCAGCTCAGAGAGACATTTCTGCTTGGGCCAATCTGGCCTTCAAGCTCAGTCCCTTCAATGAATAA
AACAAAACAAAACAAGTCGGGATTTTGTACCTTCAGTAACCTTATTGACGATTGGGAGAAAGGGAAAATGCACGGGGTTG
GAGTTACCCTTTAGACCAAGCTGACTCCTTTCTCTTATATGCACACACACACACACACACATCTTCAATAACC
TTATTGATGATTGGGAGAAAGGAAAAATGTACAAATTGGAGTTACCCTTTAGACCAAGCTGACTCCTTTCTCTTACACA
CACATACACACACACACACACACATTTCTCTCTCTCTCTCACTCTTTTCATGCCCTTACATACATGCACACACAG
AGACCAAGCTGACTCCTTTCTCTTACACACACACACATGTCATGTGCACAAACACACATTTCTCTCGCTCTCTCATGCGCT
TACACACATGCACACACATACATATTCCTCTCTAGCAACTGGCATATTCCTCCCCTTTCTGTGTAGATGAGGCACAG
ATTAGTTCCACCCAAACCAAGCTCTTCAAAGTCTCACCTTCTGTCATAAATAGCTTTATCGAACATTTTAATGCAGGC
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TGGTCTATAAGTTATTTAATTTTATTCTTAATATGGTCAAAGATAATGGTTGCACTGACTTCAACTTTACTTTCTTMAA
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TGTGACTCCTACTCTTGAAGATATAGTCTTACCTGCAGATGCCTGACATGGCCAGTCTTTGAGATGGCCAGTGGCTGAG
GATTTCTCAGATTTCTCCAGAATCTGTTCCCTTAAGCAAGGCCCTACTCATGGTTATTTCTTTTTTTGACACACACATCT
TTTTTTTTTTTTTTTTTTTTTTTTTTTGGAGACAGAGTTTTTATGCTCTTGTGTTGCCAGGCTGGAGTGCAATGGCGCAACCTCT
GCTCACTGCAACCTCTATCTCCCAGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCAAGTAGCTGGGAATATAGGCATGT
GCCACCAAGCCCGACTAATTTTGTATTTTTTAGTAGAGACAGGGTTTCTCCATGTGGATCAGGCTGGTCTTGAACCTCCCG
ACCTCAGGTGATCCACCCACCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCTGGCCGACACAGA
CATCTTGTTTCATCATAGGTCTGTGTGCTTGCTCATCTTTTCTCTTAAAAATCCTTCCCACTCTCTTTTATTTTCAATTCTT
GAGGACTCAGCCTTTTCCCACTCCACCTCTGTCCCTGTCTGATTAAATAGTCTCTTTTACTGTCCCATAGCCACAT
GCTTCCTTCTGTGAGAGTGCACTCTACAGTTGTTTGCTGAAATTACCTCCTCTGCTGGACTCTGGGTAAGGGACACACT

Fig. 9.329

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CTTTTCATCATCTTGTCTTCTGCAGCCTCTAATAAAGTGCATGGCACATAGTCAGTGCTAAATAAATGTTGAGTTACTGG
TGGGACTAAAAGTCAATGAAAGCCAACCTCATGTTTATTTTCATATAAAAATTCTACTAGAGGCATAGGCAACATTCGGA
AAAACAATTGTAGTTAGTGAGAAGATAAAAGAAAAAGAAAACCGTCACAAAATTGCACACATCTTTCCTTTGGAAGCTT
TATGAAGTACTAAATAAAGTTTTATATATTTTATACAATTTTAAATACTTCAAAACAATAATTTGATGCCAGAAATAC
ACTTGGAGATGAGAGCAGCTTGCCACTAGCAAACCTCTGCTTAAACCTATTACATGTACACATTGAAAGAGAATCCAAAG
CCTTCATGTATTTCCCATCAGATAAAATGTATAGAGGAAAAAAATTAAGTCAGCAAAAGTTAGACCTAACCTACACAA
ATCTTTTACTGTAGCAAACCTAAAGGAATGACTAGCTCAAAGCAATACACGGTGAAACAGAAATCATTTTTTCCAGTTCT
ATCTACTGTAGACAGTATCAATTCCTTCCCTAGAACAAAGGGGAAATTTTGTAGAATTAAGAGAAGAGAAGCTGGAAC
TGGTTAGGGAGATTTAAGTATTTGCTCTTAGGAGCTTTTTGTTGTAGTTCTTTTATTTTAAAAAATCTGGATCAGTGC
TCATCATGACTGGCCATCAGAGAAATGCAAATCAAACCACAATGAGATACCATCTCACACCAGTTAGAATGGCAATCA
TTAAAAAGTCAGGAAACAACAGGTGCTGGAGAGGATGTGAAGAAATAGGAACACTTTTACACTGTTGGTGGGACTGTAA
ACTAGTTCAACCATTGTGGAAGACAGTGTGGCGATTCTCAAGGGTCTAGAACTAGAAATACCATTGACCCAGCCATC
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TATTCACAATAGCAAAGACTTGGAACCAACCCAAATGTCCATCAATGATAGACTGGATTAAGAAAATGTGGCACATATA
CACCATGGAATACTATGCAGCCATRAAAAAGGATGAGTTCATGTCTTTGTAGGAACATGGATGAAATTGGAAACCATC
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CACATGGACATAGGAAGGGGAACATCACACACCGGGGCTGCTGTGGGGTGGGGGGAAGGGGGAGGGATAGCATTAGGA
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GGTTCACGCCATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCGCCCGCCACCATGCACGGCTAGTATTTT
GTATTTTTAGTAGAGCCAAGGTTTCACTGTGTTAGCCAGGATGGTCTTGATCTCTTGACTTCATGATCTGCCCCGCTTG
GCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACTGTGCCCGACCTCAGCTGTTATAATTTGTATTTTTTTTTTAA
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ATCCCGTACAGTGGATCCTTTTACCTCAATGGTCAACATCACTGTGAGTTGTGGGTAAGAGTAATGGACTGAACATTTT
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GTTCAAAGGTGCCTTGAAGAAAACCTGATGCTGTTGTAGCAGGGGCTTCAAGGCTCAATTTATTTCTCAGTTGCTGTTT
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GCCAAAACCTTCTGAAATTATTTTCCACATTAAAAAATAACAACCTGAAATATAATGTGTGGAGCCACATCCTGTTAGAT
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TGCATATGGAAGTTATTTCAGTAAATGAAAATGATTGTGTTAGAGCCATAGGCGAAAGTATTGTTTGTGATCTTTAGGG
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TTAGAATTGGGGAAGCTATACAGAGCAGGTATTTTATATTTACCATTTAAAATTATTAATATCTTTAAGCTTGTTAAGG
GGGAAAAAAAGCTTCAATCTGAACTTTAATTCACCTGGCAAGGTTTAGGCAGTTGTTTGTCCATCTAAAGTATATAT
TTACATAGACTGCGGGGTGGAGGAAGTAGAACAAAAAGAGGCAAATTTTAACCTAGAGTACTCAAAGTGAACAGTAAA
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CTCCTCACTATGAATGGTTTGCTTATTGCATGCTAATGTAGAAAAGTTCTCCACAACCTGATCTGCTTAGGGACAGTGT
CCCTGGTGCCCTGGGTGAGCCTTTGTAAACAGAACAGGCTTTTCTGTATGCCTTTGAATATGGTCTTTCCGTTTTCTCA
AAATTGTAGTGTACTCTGCACATGGTGGGTAGAAACACCTTCCAGATTTTCTTCCCTTGGCCCAAGATCATCTAGGTCT
TATGCAATTCACAGGGGAAAGGAGTAGTATGCAGGACATGCGTAAATTTCTCCTTACCATGTGGTTCTTTTACATTGCTG
TTCTTTTAAAGAGAACTCTTAGGCAATTCGGCATATGAAAATGCAGCCTATTGTCAAATTTGTGAATTATAAAGCGTTCC
GACCCCAACATTGTATATTTTGTGTTGCTTGTCTGTCGCCCCCAACATTTAGAAGTCTGGTACTTAGTAGGGACTG

Fig. 9.330

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GGCTGGGTAGAAAAAGAATGCGGTCTTAATTTCCACAACCTGCCTCCTTGAAGTAGGTGAATCCTAAGCCTTTAGAAA
CGAACCCCCAGAAACCTTAAACCTCAGAAACCTTGAGGGTCCTAAATAACCTTGTTTGTGAAACTTCGCAG
GAAGCAGCCGGTCTGAGGAGTCGCAGAGGCTCCAGCGGTCTGGGCAACCTGGAGGGCAGTGGCTCGGCGGGGCGCGCGGT
GGTTCCTAGCGAGACAGGTGCGCGGCCGTGCGCGCCTGCGTGCCGGTCCCTCCTGAGCCCCGCGCGGGGCGCGACCCCGG
ACGGCGGACTGCAGGCGCTCCACGCCGGGTTTTCCGCCGCTCTCCTAGCACAGACAGTCCCACAGCAGCCCCACAGACA
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GGTGGCGGGGCCCCCGGGGAGAGCTCGAGGGGAAGGACGCGGCGGGTGGCACGGACAGGGACAGGCTTTTGCAATTCG
GAATCTTTTCGTAAGGGGGTTGAGGAGGAGCCAGGCAGCGCCGAGGGCCGAGAGGGGCGTGAGGGGGAGTGTTCCCGGA
AAGCTGCGGCGTTGGAAGCCGCGCTGCGCCTGCCGCCGAGGAGTTGAGAAAGAAAAACAGGAAACGTGGTGGCCGCGCA
CCCGGCCGCGGCTGATTCATTCACTTCAAGTGCCGTGCAGAAGGCTCGGCAGGCGGGGCGGGCGTGGGGCCGCGGCTCC
GGGTGGGGACCGAGGAGATCCGGCTGTGGACCAGACGCTCCTCTGCGGGGCGGGCACCCAAGCGCGCTCGCCACCCCC
TCGCCATCCGCTAGAGCCGGGCTCCTGGACTGGGACTCGGGCCCCGCCGACAGTTGAAAAGTCGCATAGTGGTTTTTC
GCTCGCGTCTGTGTGAAAGTTGGCTCGCCGCTCTTTGCACGCCCTCCCTGGAGGCCGACCCGAGACGCCAAGCTGGA
GAGACCGTGCTCCCCGAGGCCGCGCCCCGCGAGCACAGCCTCCGCCCCCGTTGCACTGCCGGGCTGGGCAATATGA
AGGAGCAGCCCTCATGTGCCGGCACCGGGCATCCGAGCATGGCGGGGTATGGCAGGATGGCCCCCTTTGAACTCGCTAG
CGGACCCGTGAAGCGCTTGAGAACTGAGTCCCCCTTTCCCTGTCTCTTCGCAGAGGAGGCCTACCAGAACTGGCCAGC
GAGACCTGGAGGAGCTGGACTGGTGTCTGGACCAGCTAGAGACCCTACAGACCAGGCACTCCGTCAGTGAGATGGCCT
CCAACAAGGTAAAGCCCCGGTTCTGCTGTCACTGGTGGCCCCAGGCTGCTGATTCCCATGCCGGCGAGCCACTGGTACCC
ATAGCCCCGGGAAATAAATTGAGAGGTTTTTTTAGGCTTACTGCATGTTGGCTATTCTGCTCCTTAGAGGGAGGTTCTTG
TTTCCTTGCTTTGCCCTCCCCCTAGTCACGCCAGATAAACATTTTCCAAAAGCAATTTGACGTGCTAAATTTAAGTATCTC
CCAAGACACAGGGTTCTTAAGTAACACTGAGCCCTTGCAGCAGAAACCCAGTAGGGTCCATGGGCATTGCATGTTTAA
GGTGTCTTGGGATTCCCGGTTGAAAATAAGAATTGGTGGATGTCTGTGAAATCACTTGAATGTCACCACCTAGGGCAC
TCAAACCTCAGAAGAGTTCATCGAAGCTTGAACCTTCACCCTAATTCATCTAGTTGTGAGGGTACCCCGCAAGAACTGAG
CCTTTATATATTAGTTGGCCATACATATCCGCAGGAAACATTAACAGAGTTAGAAGGTTCTTATGATCATTACTTTT
GTTTCCATTAATCTTGGAAGAAAGCCAGTTTTTTGAGGTCAATTTAGTACAAGGAGGCTTCAACTAGGCATCTGTGCC
ATATGTGCTAAGGTGCTGGTCTTGGCAGTTAATGGAGTTTTGAGGGCTGAGAAGTAACCTTCAGCCTGGGACAGCCTTA
AAATAGCACGGAAGGCAAGGTATAGTGTTCAGTTTTGCTGTATTTTTTGGATTTCCTGTTCTTTTTGTCTTTAGTTA
TGCAAGGAGATAATTGGAATTGTGTATGTTTTTAGTGGCTATTTGCCTGTGCATGTCGTATGCAGATGGCTGAAGGATT
CAGCCAGTAAGGACTGGTAATGTTGTGAGACAATTAGTAATAGTTGCCTCGTCAAGATATTTAAGTATTTTTTGGCCACC
TATTTACAAGGTCAAGAAGGTTATATTATCTTACAGTTTATCTATGTGCACATATCTTTAAATGAGTGATGCTTTTTTT
TTTTCTACCTTTCTATGGTTTTTATCCACCTGTTCTCATGCAGTTTTTTACAAAAGGCCACGGCATAACAGCCACTTG
ATTGTCTTATCTATTAAACAGTGCTTTTTGTAGGTAGCATTGTCAAGTGAAAAGTTGGCTTCAAATAAATAAAGGGGCTC
TGTAGACACGGCCAGCAGATACTAACCTACCCATATGCACACTGACCTACCCATGTCCACACTGATGATCAATTTTTTT
GTCATTTGTGCTCATTTCTAAATTTGGCACAGCTCCTCATCAGAATGACCAATTATTGCTCTCTTACTGGGACTTTTA
CCCCCTGCTGTRGCATTAGGCACCTTTATTCTCATCTGAGTAACAAATTTCTTAGTTTTATAAAATATAGTTTGTTTTCA
TAATTTGATTCTCAATAGCAGTAGACATATGCTGAATATGTCCAGTGTCTAACTGCTAAATGGGAGCACTTTGCCATG
GGCCTGAGTTCTTAATCTATTGTGTGGTTGATTCTGTGTAAGAAAATGAAGAGCAGAATCAAAGCCACTTAGCAATG
TGCAAGCATTAGTGATRTTTTCAAGGTGATTCGACAGTAGTTTTTTCAGCTAAATGAATTTGAGCAGCTAGTTACTTTC
CCTAAAATCCATATTCTTAATGTTGAGATCTATGTTTGGATTTAAACTGAATGTGAAATTTAATAATGTATTGTAAAT
GACTTCAGCTGTCAAGGAATTAATCTATACGTTAAGATTTAAAAATTTTTTAGGTCATAATAATTGCCATAAATGATTC
CTTTCTTTTAAAGGCACCAGATAAGACAGGGATTTAAAAAAAATCTTTATTATGAAGGCCCCAGATTAGACTTTGAG
TTAATTTACAGGTTTAATAGGAGAGGTGGTCTGTTTGGTTAAATGAAATCTAGGTAAAGTGAGAAGATAATTTTTTC
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AATTGAAAATTTATTGCTGTCACTAATTTTTTAACATTAACTTAGCTCAGCTTAAGCTGTTGCTTAACTTTTAATG
TCCAAGCCATTCTGTCTTCTTTAACTTATTTATTTCTAAATCAGTGCTAAGCTTACATGACTGTTATAGACAAAAAAG
GGTTAGCACAGAAAGAAAAAAGCTCTTGCTAAAGGTTGTAAAGTTACCCTTCTTCTTCATAACAAGGGCATGAATAA
GCCACATCCACAAAAGCTGTAACTGAGTTGGAGCAGCTGATCGGAAGGCCCTGTGAGTGTTGCCACACCTTCAGTCT
GAAGTGGAAATGTGTACAGCTTGCATCATCTCCACTCAGTTATTTGATATTTCTTGAGTATGAATGAATGAGTCAGAAA
CATTGCTTTTCAGTTGTTAGTTTTATTACCATCCATTCTTTTTACATACAGTGTTCAAACCTGATATTTTCAGTTTATT
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TCAATAAAAGGCATCTTAGCCAATTAGCAGCTTTTAAATGATGCTCTAGGAGCAACTAGCTGTATTTCTGTATTGGTA
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ACACATGTAGGATGCTTGTACCAAGTAGGGTAAGAGATTCAGATGGCATTTAATTTGAGTGATTAAATCTATGGCATT
TACCCTTATAAGCACATTAATCTGCCTGAGATTTGTACAGATTTCTTTTGGAACTCATTGCTACAATTGAGGGTAATT
TTAGTGAGGTCTCAAAGCTTTGGAGGCAGGCAAACCTGGAAGTTGAATTATGGTTTTGTTTCTTGCTAATGGAGTTACA
TTAGAGAAATGACTTCAATTTTTTTTAGCTTCAGTTTCCTTAGCATACAATGGGCACCATAAATACTATCTTGAAAAGT
CAGTGTAAGAGTTTCAAAGAATATATACAAAAGAGCTAGCTAATATAGTATCTGACAATAGTAGGCACTGTATCTGTTG
TTATTGCTATTATTTGGGTTTTTAGGTCAAACCTTCATTGGTTGCCTACGTGGCTAAAATGTCTTTTTTACCATTCCGTTGT
AATTTATACCAGACACTTCAAGTTTGATGGATTTTTATAATTCTTAGATAACTGTCTATAGCRCACATTTCTGTATCAT

Fig. 9.331

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AAAAATGCTTAAATTATATTTACAGTTTTTGTGGTATAATACAGACATGAACATTCTGAAGTTCTAATTAGAAGTTTAG
GCTTTTCTTAAATTCCAGTTACAGAATGCCAGACAGAAGAGCTAACTAAGTACTTTTCTGTCTTTACTCTAAACTGAA
AATGCAGACCTTACTAATTTTGTAGAACATGTGAAATGTAATAAGGGTTAGTGGTCAATGTGCCTTTTCAGTAAACACC
TTTTAAGCAGAATAGTTTCACCATGTTTTTGGTTACCTTTCTCTCATGGAAAGTATATTGAGGATGGGAGTCAGTAGAG
AGGAGAGCAGGACAGCGGCCTGGGCCTCAGGGCTTTCAAGGATTATGCTATGGAACCTGTAAACGTTATCCTCTTGTA
CCTAATAATGCACATTTCCACCAGGCCCTCCCTGTAAGTTTCAGTGAGGTTCAAAGGAGAAAACGAAATCATTTTGAGTTA
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CTGGCTGCTCCCCCTACTGCTTTGTAATAGTCAATGCAGAACATATAGTAGGACTTTTTGTTGATGTATTTTCTTCTGG
GCAAGAGGGGTGTTATAACAAATATAGGATCTTCATAGAAGTGGCTAAATCTTAAGATATTTCCACATTATGCAACTAC
AGTGTAACCTAACAGATATAAATGTTAACTTTTGTCTAAGAAGGAACTAAGTTAATTGGAAAAGGCATGTTAGTTTAA
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CAAACCTTTAAATTTCAAAGAAAATATCTTCTATAATTATAGAAAATCAACATTTAGATGTTTTGAGTTTCGATATCTGC
TTTTTCATCTACTCAATAAAGGTAGATTGTTGGGAAAGATTATGTAGCTTACATGTAGTACCTTAAAGTTAATATGAAAG
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AACTCCTAATTTTAAATTATTCAATAAATTAATCATTACTAGATAAATTTCTTTTTTTCAGTTACATTTTGACTTAATA
TTTGGGGTAGTAGTGGTAACTTCTGTCTGAGAGCATTATGAACTGTCTACGTTTTTCGAAAAAATTCGAAACATAAGG
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CAGGATGGTCTCGATCTCTTGACCTCGTGATCCTCCCTCCTCGGCCTCCACAGTGCTGGGATTTTGTCAATTTCTGTGT
ACATTCATATCTGCTATTATATGTTGAATGTTTTAACTTAATGTATTTTTTTATTAATCTTTAAATTTATTTTRTCAATT
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CCTTTCTACTCTCTGTTTCTGAGATCGACTTTTCTAGATTCCCCATAAAGTGAGATCATTTATTAAAGACAATATTTGT
CTTTCTGTGCCTGACTTATCTCACTTAGCATAAATGTCCTGTAGTTCCATTCCATGATGTTGTGAATGACAGAATTTCTT
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GTTAGCCAGGATGGTCTCGATCTCCTGACCTTGTGATCKGCCCGCCTCGGCCTCCCAAAGTGCTGGGATTACAGATCTG
AGCCACTGCGCCCGGCCTGTATAATCTA
TCACATTTTCTTTATGCACACTTGATGAACACTTAGATTGCTTCCATATCTTGGCTGTTGTAAATAAAGCTGAAATGAA
TATGGGAGTGCAAATATCTTTTTGATGTACCAATTTTCAGTTCCCTTGGGTGTATACCTAGAAGTAGGATTGTTGGATTT
TATGGTAGTTCTATTTTTTAGATTTTTGAGGAACCTTCATACTATTTTTCAATAATGGTGGTTCTAATTTACATTCCCACC
AACAGTGTAGAAGGGCTCCCTTTTTCTCCACATCCTTGCCAACACTTGTATCATTCATCTGTTTGAACATAGCCATTGT
ACATGGTGGAGAGCAACACATACTGGGGTCTGTTGGCGGGTGGGATGGGTGCAGGGAGAGCATCAGGAAGAATGGCTAG
TGGATGCTGGGCTTAGTGCTTAGGTGATGGGATGATACGTGAAGCAAACATGGCACATGTTTACCTATGTAAACAACT
GCACATCCTGCACATCCTGCACATGTACCCCTGAACTTAAAAGTTGAAGAGAAAAAAAAGGAAAATAGTCATTCTAACA
GGCATGAGGTGATAGCTCATTTGTGGTTTTAATTGCATTTCCCTGAGAATTAGAGATATTGTTGAGTTTTCTGTTTGT
TGAGACAGGGTCTTGCTCTTTTGTTCAGGCTGGAGTGCAAGTGGCGTGATCATGTCTCACTGTAACCTCAAATTTCTCGG
CTCAATGATCCTGCCTCAGCCTCCCAAGTAGCCAGGACTACAGGTGTGTGCCACCATACTGGCTTTTTTATTTTTTCTG

Fig. 9.332

ATAGAGAGGGTCTTGCTATTTTGGCCAGGCTTGTCTAGAATTCCTGGCCTCAAGTGATCTCCTGCCTCAGCCTCCCAA
GCACTGGGATTMCRRGCATGAGCCACTGTGCAGGGCTGGGGAACATTGTTTTTTTTTTTTTTTTTTTTTTTTTTGTATCT
CTGTTGGCCATTCATATTTCTTTTTTGGAGAAAGTGTGTCTATTCAGATCCTTTGTCTAATTTAATCGATTTGTTTTCT
TTACTATTTGGTTGTTTGAATTTTTTATATATTTTGAATATTAGCCTCTTATCAGATGTATGGTTTGCAGATATTTTCT
CCTGATCCATGGGTTGTCTTTTCACTCTATTATTTGGTTGCTTGTGCAGGTACTTTTTAGTTTAATGTAGTCCATTTG
TCTATTTTTGTTTTAGTTGCCTGTGCTTTTGGAGTCCATCCAAGAAATCATTGCCAGACCATTGTTGTGGAGATTTT
CCCTTATATTTTCTTCTAGTAGCTTTACAGTTTCAGGTCTTATGTTTAAGCCTTTATATCTCTTTTGAGTTGATTTT
TATGGAGTGTGAGATAAGAGTACAGTTTCTTTCTTCTGCAGGTGGACATCCAGTTTTCCCAACACTATTTGAAGAGACT
GTCCTCTCTCCATTGTGTGTTCTTGGACCCTTTGTCAAAAATCAATTGACTGTAAATACTTGGATTTACTTCTGGGCTT
TCTATCCTGTTCCATTGGTTGATGTTTGTTTTTTATGTCAATAACATGCTGTTTTTGATTACAGTAGCTTTATAATATATT
TTGAAATGAGGGAATATGATGCCTGCAGCTTTGTTCTTTTTTGTTCAGATTGCTTTGGCTATTAGGTCTTTTTTTGGTT
CATATAAATTTTAAGATTTTTTCTCTATTTCTGTGAAAAAAGACATTGGAATTTTGAAAGGAATTGCATTGAAT
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GAACATGGGATATCTTTTCATTTATTTCTGTCTTCTTCAGTTGTTTTAAATCAATGTCTTGTAGTTTCCAGTGTATAGAT
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CAGGCTAGAGTGCAGTGGCGCGATCTCGGCTTACTGCAAGCTCCACCTCCAGGTTACGCGCTTCTCCTGCCTCAGCC
TCCCAATTAGCTGGGACTACAGGCGCCTGCCACCACACCTGCCTTTTTTTTTTTTTTGTATTTTAAAGTAGAGAYGGGGT
TTCAGTGTGTTAGCCAGGATGGTCTCGATCTCCTGACCTCATGATCTGCCCGCCTCGGCCTCCCAAAGTGCTGGGATTA
CAGGAGTGAGCCACCACGCTGGCTGCTCCTAAGTATTTTTTGTATGCTATTGCAAATGGGATTATTTTCTTAATTTCT
TTTTTGGATAAATTTATTGTCAAAGTATAGAAATGCTAAGAGCAACTTATTGTTAAATCTAAATACTCACCCAAGTGCCT
CATCTTAAGTAATGGTATACATGAAATCATAGGTTTGATGTTCAAGTTATATTTTTTCTGTATTTTTTCTAAAATTAAT
GAATAACAAAATGAGAAAACATTTCATCTTGCACCACCTTAAAGCATTTTGCCTACCATTACTAACACTGGGCATACTT
TTGGACACACTAGTCTAAATCAGCTGCAAGAACAATTTAGAGACTAATTTATTGTTTTTAAACAATCAACATTTTTCTT
CTTTACCCTCAGTTGGATTTCTCCACTAGAGGAGAAAATGGCAAATTCGTTTCAGCAGAAATATTCCAATTCATAAAATA
TTCATTCTTAGGCAGAATGGTCCGTGGATTAGAGCACCTACTATAGACAGCCTGCCTGAATTGGATTCAATCTCTG
CCTTTTACCAAACGTAAATTGGGCCAYGTTGCCTTCTGTGCTTCAGTTTTCTCATCTGTTAAATAGTTGCTGTGGTGAT
TTAATGAATTAAAGATGGGAATTGCTCTCGACTGAGTCAGGAATCATACTAGATACTCAGTAGAACTCATTCATTTCCA
TATGTGAAGGAATAACATTAGTTTTTATTATTATTGTTGGGGGTGAGGGTTAGAGAAGGGTGTCTGCCATCCTGATT
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CCCCATTAATCACTTTTTTTGGTGGTGTTTTATATGTTTTATTCTTTATTTTCATCCTTCAAGGAGTTTGCCCTTTGT
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TTCTAGCAGAAATGCATAATGCAGTGCAGAAGAATAGTCTAGCCAGTGTTTTTTCATAAATGAATGTACATACGTTCTTC
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TAGAAGCTACCATACCCCAGTAGCAATGAGCACACTTAGCACCTCATTTTGGTTTCTAATACCATTCTCTGGTAAAAG
GAACCAAGACTCTTTGAAAAAAATAGCTGATTTCTAGGGACTCAGGAAGGAAATATGTAAGATGAGCCTGGAGCATCT
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GGTAAATAAATATCCATAAACCCATATTGATACAATTTATTAAATAAATACCTAAGTGGGGGAAAATAGACAAATCTC
CTCCTGTGCGGAATTCCCAATAACTTCTATAGATACTCTACTCTCAAGGAGGGAGAACATAACCCCTACACCGTAAGT
GTGAGCTGTGCATATTGACTTCTTCTAAACAGTACAGTGTAGAAAGGGGAAAAAAGAACTGTAGTGGAGAAWCCTA
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CAAGGAAAACCTGAGATACCTACAGCCAAGAAGAGCCTCAGGAGACATGAGGACTAAATATTATGTGGTATCCTGGATG
GGATCCAGAACAGAAAAAGGAAATTACGTGAAAATAAGGAAATCTGAATGAAGTGTGGACTTTGTTTAAATAAATG
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CCTATATCTGGGGTGTCTATGTCTCCACCTCAGGCTTTCTAATTGGATACCTTTGACTTTCCACAGAACTTTTATT
TGTAATTCTCTTAACTCTTAGTACAATATTTTGTGTCTCTGTTTTATTGTTCTGATGAGGTCTTCTACGGTGTTTTTAG
CCCAACCCAGTGAGTGTCTGGTGCCTTAGACCCTTGGCCATCCTGACACCCTACACAGTGAGTGTCAATATAACATTT
TGTAAGTGTGACATAAATTAGTACAAGAATGCATGATGTTGGTTAATGCAAATGCTGTTTTTTTACCTTGTATTGGTAG
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ATTTCTCACTTAATCTCAGTTCTGCTTTTAGATACTCATGTCTCAGAAATTTTATCATAGTATATGCTAGATCAATCTC
ATCTCGTTTAAATTAAAAGTTTGTAGTGCACAGGCAAAAACCATAGAGGCCATATTCATGGTCATAAACTGCTCTTTCT
CATCTCTACCTATTTTAGCAATAACATTTCTTCTTATTAGTTTGTTCAGGTGAAAATGACATAACTGAATACTTTCCCT
GCTAGTCTGGGAAGAAAGGATGTGTTTGTGGCTTTTGCCTGTGAGAGTATTCTGCAATGTTTGCCTAGGGCATGCT
TGCAATCCCATCAGCATCTCTGGTCTGCTGCACTGTAATCCCTATAGGGACAGCCTCTGGCTTTTATTACTGACAAGCA
GTGCACTGTGCAGCCATAGGCACCATAATAGGAAACACCTTGGCCTGTGATAAACAGGTCTGGAGAGTAGAAAGTACAG
GCCTGCTGGGGATGTGTCCATAGCAAAGAGGCAAAGATGCGGCTGCCATATTGGAGTAAGTGACAGGCTAATGTCTGCCA

Fig. 9.333

TCTCCATTTTCAGAAAATAACTGGCTGATTTTGAAGCTGCTTTTTGTATAAACAGTAGTGTTTTGGTTGCTTTTTGT
TGGCTTAAATATGAATAAAGCCATCTTAAAGAGATTATACCCTTCAAAGTATTTTGAGAAGATCTATAAAGTATTTTCC
TTTTGTTATTTTACATTTAATTCTACCTGATCATTTCCAATCCAAACCCAATAGAGAAGGAAAAACAGATATTTCACTAT
AGTGGGAAATTAGGAAAAAAGAACCATGCAAAAATACAAGTGATTGTGTGTCTTTTAAAAGAATTACAAATCACACTG
AATTACCCAAAATTACAAAGAAAAGTGCATTTATTATTAAAGGTAACCTTGTGTTGTCTGTGCCTTTACATCAACTCCAAG
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AACTGCAATATTAATGATAGGCTTTGTAGCAAGAATTTAGGAAGACAATAAATTTCAAATTTGGAAGGGTTATCACAGT
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AGAGAATGCATCATTACAGACTGAATCATCAGTCCCTACAGAAGGGGAATTTGTTCTTTCAAAGTAGAATTTTCAGCAG
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TCTCAGCTACTCGGGAGGCTGAGCAGGAGAATCACTTGAACCCAGGAGGCAGAGGTTGCAGTGAGCCAAGATCGTGCCA
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GGCTCACACCTGTAATCCCAGCACTTTGGGAGGTGATCACATGAGGCCAGAAGTTTGAGACCAGCCTGGCCAACATGG
CAAAACCTGGCCTGTACTAAAAATACAAAAATTAGCCAGGTGTGGTGGTGCATGTCTGTTATCTCAGCTACTCGGGAGG
CTGAGCAGGAGAATCACTTGAACCCAGGAGGTAGAGGTTGCAGTGAATGGAGATGGCGCCACTGCACTCCAGTCTGGGC
GACTGAGTGAGACTCCATCTCAAAAAAAAAAAAAAAAAAAGTAGAATTCATGGGAAAAATTATTTGTTGCTATTTAGCC
TTATTTTTTAAATATTTTCAAGTATATTGTCTGTTTGATACATATGAACAATCTGACTACAACCTATTGGGAAACACCAGTA
TTTACCTTACCTTCTAATGTAAGGCATGATTCAGGTATTTTCTCATACTCAACCTTAAATCTCTAATTTAGTCCCA
GAAACAGTATTCTACATGTCAAACGTTTTTTTGTTTTGTTTTGTTTTTGGAGACAAGTTCTCACCTTGTCAACCAGGCT
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GTGGATGGAGCCACAGGCATGTGCCACCAGACCGGCTTTTCTGTTTTTTAATTTAATTTAATTTTATTTTATTTT
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CGTAAGCCACTGTGCCCGACCTCTGGTTGTATTTTCAAAATATATCCAAATAATTTTTAAAAATGTATTTAGTGAGCA
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ATGTGCACAGAAGGATAATAGGGTCAGTGGGATGAGGAGACCAAGTGACTAAAGCAGATSTGAGAATCTGAGCTGTAAA
GGATTTGGAGGTGGGATGGAGGGATAAAACCTTAGTGGAAGCAGACAGCAGATAGAGAAGATATAGGATAAACTCACG
GAAGGTGTAAGCAGGTTTCATGATTTTAAAATTAGCCTRTTAAGAGGTTGTGTTGGAGGTCTGTTTTCTTAGTGGGGGCA
AAGTGGCAAGGGAAGTTTCACTTCTGAATAGGGTGGAATGAAGATATAAGGAAAGAAAGAGAGAAAGAGGAAAGACG
TATTAGGCAGTGTGAGGCAGAGAAAGAGAAGAGTGGGAGAAGGAGGGGAGTCAGACTTTTCTTGGCTGGAGGCAAACCTG
ACCAATGCCCGTTGCTGCCCCCTTTCCACCCGAGGGTTTCAATTTGGTGAAACCTGGTTGGACTGAACCTTTAGGAGAACCT
TGAAGTAAAAAAGAAATCCCAGACAGGTCAGTGGCGGATGTAAATGTTTAAGGAAGAACAGACAGTCAGCTGTAGAAAT
GGAGGGAGGGGTAGAGCTGACTGGAGAGCGATTATGGATGGTATATTTTCATGAGGCTTGAGTCTTTTTTAGCACTAAT
AGTTGGCGTCAGCTTCACCTGTTGCCCTCCATGAGTCATGCCCTGGTAGCAGAGGATGGTGGGCCAACTGCCAAGCCATC
CCCTGAAGGACCGGCTGCCCTGGGAGTGAGGAAAGGTGGTTGATGGGCTCTGAGACAGCAAGGTACAACTGAAATGGGG
TGAAAAGAGACTGTCAAATAAGTATGGGCTGATTTGTTCTAATATATCATAGGTTATTATTAGATGCTGGAAGAGTAA
AATGGAATAGAAGATGAAAAATGTGAACCTTTATCTTGATTCCATTTTAAATCTCCTAAATTTCTGAGGAGCTTTGCAAT
TCCTCTTCTGGATAATAAATCTGCCCTACAGTAGAGTTGCAACCTTGGATAAATAAATAAATTTGGCCTCCCTGAGCCTC
TGTTTTCCCTGTAAATCAGGACTAACATAATCTACCTCAAAGGATTGTGTTGAAAATTAAGTGAAAACCTTACATAAATT
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AAAAGGTCAACTTTATATGCTGCCATATAAGTAACCCAGTGCAATTGGAGCTTGAATCAACAGGGCTGGGAGAAAGCTT
ACAGAGTGTGCCCTTTGAAATCCCAATACCCAGTCATGGTGATTTATGAGACCTTACCAGCTTGCATATGTGATGGCCCA
CTGAGTTTGTTTACTTTAGTTTTAGCTCAGTTACCCAGCACTCTATGATAGCCTATTTTTTTATTACACTCTCACTAAC
TCCTTTATTATGATTTCTTTGAAAGAATATGAAGACAGCCATAAAAAAGACTCTTATTGAAGTTGAGAGGGTCTCCTGT
AGGTCTTCACTTCACTGTGTCTAAACATTGGTCTGTTGCTTGTGCTTTTCTCTCACCCCTGGATGACCTTCAAGG
CTCTTCCAAATGATTTCTCTGATTCCCTGAAATACTGAAATTGTGATTAACAGAAGTTTCAAGTGTTAATCAATATCAA
GATAGTAGTTGTCTCTTCTGGGGAAACTAGAAATGAAAACCAACATAGGCCTCTGTAATCATTCACAGATTAAAAA

Fig. 9.334

Fig. 9.335

AGTGTGTGATGTTCCCCCATCCTGTGTCCAAGTGTTCATTGTTCAATTCCCACCTATGAGTGAGAACATGCGGTGTT
TGGTTTTCTGTCTTGCAACAGTTTGCTCAGAATGATGGCTTCCAGTTTCATCCATGTCCCTACAAAGGACATGATGAA
CTCATCCTTTTTTATGGCTGCATAGTATTCATGGTGTGTATGTGCCACATTTTCTTAATCCAGTCTATCATTGGTGGA
CATTGGGGTTGGTTCCAAGTCTTTGCTATCCTGAATAGTGCTGCAGTAAACATACATGTGAATGTGTCTTTATAGCAGC
ATGATTTATAATCCTTTGGATATATATCCAGTAATGGGATGGCTGGGTCAAATGGTATTTCTAGTTCTAGACCTTTGAG
GAATTGCCCACTGTCTTCCACAATGGTTGAACTACTTTACAGTCCCACCAACAGTGTAAGAGTGTTCCTATTTCTTCA
CATCCTCTCCAGCACCTGTTGTTTCTGACTTTTAAATGATTGCCATTCTAACTGGTGTGAGATGGTATCTCATTGTGG
TTTTGATTTGCATTTCTCTGATGGCCAGTCATGATGAGCATTTTTTTCACGTGTCTGTTGGCTGCATAAATGTCTTCTTT
TGAGAAGTGTCTGTTTCATATACTTCACCCACTTTTTGATGGGGTTGTTTGATTTTTTCTTGTAATTTGTTTAAAGTTCT
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CTCTGATGGTAGTTTCTTTTGTCTGTGCAGAAGCTCTTTAGTTTAAATTAGATCCCATTTGTCAATTTTGGCTCTTGTTC
CATTGCTTTTGGTGTTCAGACATGAAGTCCTTGCCCATGCCTATGTCCTGAATGGTATTGCTTAGGTTTCTTCTAGG
GTTTTTATGGTTTTAGGACTAACATGTAAGTCTTTAATCCGTCTTGAATTAATTTTTGTATAAAGTGTAAGGAAGGGAT
CCAGTTTCAGCTTTCTACATATGGCTAGCCAGTTTCCCAGCACCATTTATTAATAGGGAATCCTTTCCCCATTTCTT
GTTTTTGTGAGGTTTGTCAAAGATCAGATGGTTGTAGACATGTGGTATTATTTCTGAGGGCTCTATTCTGTTCCATTGG
CCTATATCTCTGTTCTTGTACCAGTACCATGCTATTTTGGTTACTGTAGCCTTGTAGTATAGTTTGAAGTCAGGTAGCA
TGATGCCTCCAGCTTTGTTCTTTTTGCTTAGGATTGTCTTGGCAATGCGGGCTCTTTTTTGGTTCCATATGAACTTTAA
AGTAGTTTTTCCAATTCTGTGAAGAAAGTCATTGGTAGATTGATGGGGATGGCATTGAATCTATAAATTACCTTGGGCA
GTATGGCCATTTTCATGATATTGATTCTTCCTATCTATAAGCTTTGTGTCTCTTTTTATTTTGTGAGCAGTGGTTTGT
AATTCTCCCTGAAAAGGTCCTTCACATCCCTTGTAAGTTGGATTCTTGGTATTTTATTCTCTTTGAAGTAATTGTGAA
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CATGTCATCTGCAAACAGGGACAATTTGACTTCCTCTTTTCTAATTGAATACCTTTGTTTCTTCTCCTGCCTGATT
GCCCTGGCCAGAACTTTCAACACTATGTTGAATAGGAGCGGTGAGAGAAGGCATCCCTGTCTTGTGCCAGTGTTCAAAG
GGAATGCTTCCAGTTTTTGCCCATTCAGTATGATATTGRCCTGTGGGTTTGTGATAAATAGCTCTTACTATTTTGAGATA
CATCCCATCAATACCGAATTTAATTGAGAGTTTTTAGCATGAAGTCCTGTTGAATTTTGTCAAAGGCCCTTTTCTGCATCT
ATTGAGATAATCATGTGTTTGTCTTTGGTTCTGTTTATATGATGGATTACGTTTATTGATTTGCATATGTTGAAGC
AGCCTTGCATCCCAGGGATGAAGCCCACTTGATTAGGGTGGACAAGCTTTTTGATGTGCTGCTGGATTGGTTTGGCAG
TATTTTATTAAGGATTTTGTGATCGATGTTTCATCATGGATGTTGGTCTAAAATTCTCTTTTTTGTGTGTCTCCGCC
AGGCATTGGTATCAGGATGATGCTGGCCTCATCAAATGAGTTAGGGAGGATTCCCTCTTTTCTATTGATTGGAATAGT
TTCAGAAGGAATGTTACCAACTCCTCTTTGTACCTCTGGTAGAATTCAGCTGTGAATCTGTCTGGTCTGACTTTTTT
TGGTTGGTAGGCTCTTAATTATTGCCTTAATTTCAGAACCTGTTATTGGTCTATTACAGGGATTCAACTTCTTCTGATT
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TCTCTGATGGTAGTTTGTATCTCTGGGGGATTGGTGGTGGTATCCCTTTATCATTTTTTTATTGCATCTATTGATTCT
TCTCTCATTCTTCTTTATTAGTCTTGTAGTGGTCTATCAATTTTGTGATCTTTTCAAAAACCAGCTCCTGGACTC
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AGCTTTTGAATGTATTTGCTCTTGTCTTTCTAGTTCTGTTTAAATTGTGATGTTAGGGTGTCAATTTTAGATCTTCTCTGC
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GCAGGTTGTTTCAGTTTCCATGTAGTTGAGTGGTTTTGAATGAGTTTCTTAATCCCAACTTCTACTTTGCACTGTGGTCT
GAGAGAAAATTTGTTATAATTTCTGTTCTATTACATTTGCTGAGGAGTGCTTTACTTCCAACCTATGTGGTTCAGTTTGG
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CGCTTGTGTCAGAGCTGAGTTCAATTCCTGGATATCCTTGTTAATTTTCTGTCTCGTTGATCTGTCTAATGTTGACAGT
GGGGTGTAAAGTCTCCCATTTATTATTGTGTAGAAGTCTAAGTCTCTTAGTAGGTCTCTAAGGACTTGCTTTATGAATC
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ATGGCCTTCTTTGTCTCTTCTGATCTTTGTTGGTTTAAAGTCTGTTTTATCAGAGACTAGGATTGCAACCCCTGCTTTT
TTTTGTTTTCTATTTGCTTGGTAGATCTTCTCCATCCCTTTATTTTGGAGCTATGTGTCTCTCTGCATGTGAGATGGG
TCTCCTGAATACAGCGCACTGATGGGTCTTGAATCTTTATCCAATTTGCTAGTCTGTGTTTTTAAATTGGAACATTTAG
CCCATTTACATATAAGGTTAATATTGTTATGTGGGAATTTGATCCTGTCTTTATGATGTTAGCTGGTTATTTTGGCCAT
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AAGAATGTTGAATATTGGCCCCCACTCTCTTCTGGCTTATAGAGTTTCTGCTGAGAGATCAGCTGTAAGTCTGATGGGC
TTCCCTTTGTGGGTAAACCCGACCTTTCTCTCTGGCTGCCCTTAACATTTTTTCTTCACTTTGGTGAATCTGA
CAATTATGTGTCTTGGAGTTGCTCTTCTTGGAGGATCTTCTGTTGGCATTCTCTGATTTCTGGAATTTGAATGTTGGC
CTGCCTTGTAGGTTGGGGAATTTCTCCTGGATAATATCCTGCAGAGTGTTTTCCAACCTGGTTCCATTCTCCCATCAC
TTTCAGGTACACCAATCAGATGTAGATTTGGTCTTTTTCACATAGTCCCATATTTCTTGGAGGCTTTGTTCAATTTATTTT
TACTCTTTTTTCTCTAACTTCTCTTCTTGGCTTCAATTTCAATTTCAATTTCAATTTCAATTTCAATTTCAATTTCAATTT
TCTTCCAGTTGATGGAATTGGCTACTGAACTTGTGAATGCATCATGTAGTTCTCATGCCATGGTTTTTCAGCTCCATCA
GGTCATTTAAGGTCTTCTCTATGCTGGTTATTCTAGTTAGCCATTTGTCTAATCTTTTTTCAAGGTTTTTAGCTTCTTT
GCGATGGGTTTGAACATCCTCCTTTAGCTCGGAAAAGTTTATTACCCATCGTCTGAAGCCTTCTTCTCTCAGCTTGTCA

Fig. 9.337

ACCTCCCACCTCCCACCTCAGTAGTATCTGGAAC TACAAGCACATGCCACCACACCTGGCTAATTTTTAAATTTTTTGT
AGAGACGGTGGTTTTCTCTATGTTGCCCAGGCTGGTCTCGACCTCCTGGCCTCAAGCAATCCTCCCGCATTGGCCTCCCA
AAATGTTGGGATTACAGGCATGAGGCACCTTCACCAGGCCAAGAACTATTTTTCTATTTCAACAAACACACTTGCATA
TATGTATATAAAATCATTTGGGCATTTGAGAGCAGTGGATTACAGATAAGAAACCTGAGCTCTAGCTGTAACGCTGTCC
CTCAAGTTGTGTTGTGTCAGAACTTTTCTGGACCTCAGTTCCTTGTCTGAATGTGTGTCGTCATCATTACATCTGCATATGAG
GGCAGCTGTGTGTTGTTTCACATAGCGCTCCCAAACATAAGCGTGTCTCACTATATGGCAGGGCTGTCTGCCTGTGGGCA
CCTGCTTCCTCACCCTGTCCAGAGATCTGACCATGGTGATAGTAACCATGATTCTTTAATTCAAGGCACTGTAAAGTTA
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TGTAAGAAGGTCCATCTTTTCATGTTAAATAGGGATAATATTTATCTTATCAGAATGTTGCCAAGATTAGAAATGAGG
FATGTAAAGTTCCTTTGTGCATAGTAGGTGCCTAGTAAATGTTGTAACCTATTAAAGTTTCTTCATTAAATTTGGTGAAG
CCAAGTCTGACTATAAGAATTGTATCTCTCTGGCTCTATTTCAAATTTCTCTTCTAAATTATCTAGATTCTCTCTGCAGA
TAGCAGCTACCGTGGCAATAGGAAGGAGATTCTAGTCTCCTAGAAATGGAGATTAGGGAAATGAAATGAATTTTAATT
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TAAATAATCTCCCTTATTGTCAGCAAATTAGGGACTTATTTGAATAAGTTAAATCCTTTCACATCCAGCACCTATTAGAA
TGCCTGGCACATAATAGTTGCCAAATAGAAGTCTGTTGAATGAAGGGACCATCACTCACATTCAGCAGGGAGAAGAGGC
TGCAGATTTTAGAGGGGAAGGGAACTGTATGTGTGTTTCTGCATAATGTTTTAAGACAAGGAGTATTATCTACTATATG
TAATCTGTTTTAAATGTTTTTGATGATTTTGCTGAGGGTGAAAACCTTGTCCTTTCTGTCACCTATAATCAGTATAA
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CAGAGGCCAGGGAGGCTAGATGTCCTGGGATACATGGGACAGCCTTTCCACCAGGGAGGTTTGCTGTGTGTCCACACAA
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ACCCTTAGATAGCACTTACTGTTCTAAGAGCCTTATGTATATGCACTAATTTAATCCTCACAACAACCCTGTAAGGTAG
AAACTATTATTCTCATTTTCCATGTGAGTAAACTGAAGTATGGGAAATTTAAATAGCTTCCCCAAGGTCACACAGCTAA
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CACCAAGTATGTTTTGCCTCGAGGCCTTTGCTTTTGCCATTCTCACTTGGTTTTTCCCTTCTCTACCCTACCTCTCACC
CCGCAATCCACATGGCTTGTTCCTTTCTTCAGATCTCCTGGTATACGTCACTCATCAGAGGGGCTTTCCATGATCAC
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CAAATGTAATAGAGCAGCTGAAAATAGGCTTGGCGTTGGGGATTAAAGGTGAGATATGGCAGTCTGGAATTTACAGAGGG
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ATGGAGTGCTGTGATGCAATTGAAAGGAAATATAACATATATATTTTAAAAACAAATGACAAACCAGAGAATTTTTTTA
AATATTAGTATAATAGAATGATTATCATAATGATCATTGTCACTGTCTATCTTTACTGTCTGATACCTACTTATCTTCAA
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CTGGCTGGATGGCTAGGATGGCTTCTTGCCCCCTCAGGTCTGGCTCTGGGCTGGGGTGGCTGGCATTGCTGAAGACTGGC
CAGGCATGTGTGCTCTTTCTTGTGCAAGTGCTCTCTTGGCGCTTTCTCTCTTCACTGTCTGTGAAGTTAACATGGACTTC
TTGTAGCATAGCCTCTCAGAGTAGTCAGACCTTTTTTTTTTTTCAACCTAATGGCTTAGGGGTCCCAATAGCACATTCAA
AGAAAGCAGGATGAGAGCTGCAAGATTTCTTATGACTTAGCCTCAGCAGTCATGCACCATCACTTGTTGGTTACACAAG
CCAATCAGATTCAGTGTAGGACAGTGTGGCTCATTAAGAGTATCTTTGGAGACTAGCTGCCACAGTAATCCACTTTAC
AGATGAGGAAGCTGACACACTAAAAAGATTATTTGTGAGTGATTGTGCACAAAAGATTTAACTTGCAATTCCTGTTCT
CAGTACCAGAGTGAACCTGGGAAAATCATTTTTTCTCTCTAGGTCTCAACTTTATGTGGAAATGAGGGAGTTAGAGGGG
GTGATATTAAAGTTACTTTCTATTTCTCTCAGTCTTTTAAATATTTGGATTACAGATTACGGGAGGGGTAATGGTAAGGA
AYAACTGAAAGAAACAGCTCTGGATAAAAGATAGGGCTTGCTAACAAAGTTGAATGAAGGAGGGAAGGTGATGCCGTA
TAGGGGAACATGCCGGGTACTCAAGGATTCAGCTAGATGGTTTATATTAAACACCTGACAGTGTCTGACGCACATGAA
ACACCTGTTGAATGGTAGGAAAGAGTCATAGATTCCATCTGAGATACTGGAAGAAATATAGGCTCACTGACTGAAATGG
AAGTAATTATACTAAGAACTACTTTAAATAGAAAAATGTCTCACAGTTTTGAAAAAGTTTGTGCTGACTAGCAGGGCAT
TCGAGAGGAGAATCTTGTATTCAGAAATATGGCACTGAAGTTCAAGTTAGAAGTGGTCATTGAAGATTGTGGGCCTAGG

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GAAAATTAAAGAAGTTCAGGATCCTGGGAGTTGAGGGAGAGTAGTAGGAATGATCAAACATGTGGAACACTTCAGAGAA
GTAGGGAGACCAGAGAAAAGGAGGACCTTAGTATTCCTCTGGGGGTAGCTTTAGTGGAATGTTGGAGCTTTAATTTCTT
CCTTTTGTAGAGGAGTAAGAGTGATGGAAATGAATACAACAGATGTTAACAAAAGCAGAGGACTGTAGCAGCTAAAGGC
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GGATTCAAAGCATTCACTTAGAAATCCTCCCCTGTTTTTTTTAGTTGCAACCCCTAAATCTGTGTATTGTTTTTCAGACTA
CTTAGGCCAAAACAATTAGAAATTCATCAATGGAAAATATTTAGAGGTCACTTAAAAAAATAAACTAAATGCTTAAAT
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TGATGGGATTTCCAGAGAATAACATTGTATTAGCTTTAAATCAGTCCTTCCCCCTTTGGTAATTTTATGTAGTTATCT
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TTTAGTACATCTAGCTATGCACTCCAAAACCAATTTGTGAGATCAACTACCAGTTGAGAAAGCACTTATGGTAATTTTT
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AGAAGTAATTGCTTCAAATATTGTCTTTTATAATTATGTAAAATGAAATGTTGACTTCTTGGAGTCCCTTATAAGCC
TTGGTAGGGAGGTGGGCATGTGATGGAGGATTTCTCCAATCCATGTTTTTGTGTTTTTAAACAAAGGCTGGAAAGTACTC
TGGGAATAATGTATATGACCAGAAAGATGAACATGCAGGATGTCACCTATCTAGTCTGTACAATATTTAGATTCCTTTC

Fig. 9.339

Fig. 9.340

AGAACATTTTCTAAATTTCTCAAGATAAGCAACATGAAGTGGAAATTCCTTCTCCAACCTCAGAAGGAAAAGGAGAAAA
GAAAAGACCAATGTCTCAGATCAGTGGAGTCAAGAAATTGATGCACAGCTCTAGTCTGACTAATTCAGTATCCCAAGG
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ATGCCCTTTTCATGAATTTTCACTTCTAGTCTTTTAAATGGATAGAAAAGTTTATTAATTGGGGTTTTTTTTTTCTTT
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GGGGGTGATTATTTGCATCTTCATTTCAATTGTTTGGCCTTAAAGAGTTAGAAATCTGTCATGTCTTCTGGGTTTTAGGA
AAACAGTTAAGGTTTTGTGAATAGATTTTATCCATTAAAGAAGTGCTTTGTGATTGATACTTAAAAAGTACATAAATGT
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ACTTTCATGCCTTTTTTTTTTTTAAAGTAGAAAAATTTGTTTCCAAAGTGCATGTACATGCCACAACCACGGTCACACCT
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TTTTTTTACCTCCATGTCTATCCGAGCAAGGTGGACATCTTCACGAACAGCGTTTTTAAACAAGATTTTCAGCTTGGTAGAG
CTGACAAAGCAGATAAAATCTACTCCAAATTTATTTTCAAGAGAGTGTGACTCATCAGGCAGCCCAAAAGTTTATTGGAC
TTGGGGTTTCTATTCTTTTTTTTATTTGTTTGCAATATTTTTCAGAAGAAAGGCATTGCACAGAGTGAACCTAATGGACGAA
GCAACAAATATGTCAAGAACAGGACATAGCACGAATCTGTTACCAGTAGGAGGAGGATGAGCCACAGAAATTGCATAAT
TTTCTAATTTCAAGTCTTCTGATACATGACTGAATAGTGTGGTTTCAGTGAGCTGCACTGACCTCTACATTTTGTATGA
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ACCAATACAACCTAATCCTATTTGGTTTAAATGATTTTACCATGGGATTAAGAAGTATATCAGGAACATCCCTGAGAAAC
GGTTTTAAGTGTAGCAACTACTCTTCTTAATGGACAGCCACATAACGTGTAGGAAGTCTTTTATCACTTATCCTCGAT
CCATAAGCATATCTTGACAGAGGGGAACTACTTCTTTAAACACATGGAGGGGAAAGAAGATGATGCCACTGGCACCAGAGG
GTTAGTACTGTGATGCATCCTAAARTATTTATTATATTGGTAAAAATCTGGTTAAATAAAAAATTAGAGATCACTCTT
GGCTGATTTTCAGCACCAAGGAACTGTATTACAGTTTTAGAGATTAATTCCTAGTGTTTACCTGATTATAGCAGTTGGCAT

Fig. 9.344

CATGGGGCATTTAATTCTGACTTTATCCCCACGTCAGCCTTAATAAAGTCTTCTTTACCTTCTCTATGAAGACTTTAAA
GCCCCAAATAATCATTTTTTTCACATTGATATTCAAGAATTGAGATAGATAGAAGCCAAAGTGGGTATCTGACAAGTGGAAA
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AACATTCCTGTGTGCTAGAGAGTACCACCTGTCTACCAAGGGGAAAACAACCTTGTGTGAGGGGAATCATAACAGGGCT
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TATCTCCAATATAATCTGAGTATCAGGCCTCTGTGTTGTTCCAGCAGAGGTTCTTACAGTCCCTCAGCTATTAGCTTC
CTGGTCACATGTGGTGTACAGGAATATATGCTGGTTTGGGGCCCTGCTTCCACACAGTGTGTGCATCAAAGAGCCTA
TCCTTCTGGTGTTTTATTACACCTTGCTGTGGTCTGAATGTCTGTGTTGAAATGCTAACCCTAGGTGATGCTATTAGG

Fig. 9.345

AGATGGGGGCCCTTTGGGAGATAATTAGGTTATGAAGGCAGAGTCCTCATGAATGGGATCAGCGCCCTTATAAAAGAGG
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CAAGAGAGGAGAAAACAGACACAAAGTGATTTCTGGGCTTTTGACAACCTTTTTTTTTCTCAAAGTAGATTGACAATATT
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GGAACCTTTGTGAAGCATATGCAAGTATCAATAGCCAAATAGATCAAACCAAAGAAAGGATATCAGAGATTGAAGATCAA
CTTACTGAAATAAAGCATGAAGACAAGATTAGAGAAAAAAGAAGGAAAGGAAACAAAGCCTCCAAGAAATATGAGACTA
TGCGAAAAGAACAACCTACATTTGACTGGTGTACCTAAAAGTGATGGGGAGAATGGAACCAAAGTTGGAAAACACTC
TTCAGGATATTATCCAAGAGAACTTCCACAACCTAGCAAGTCAGGCCAACATTCAAATTCAGGAAATTCAGAGAACACC
ACAAAGATACTCCTTGAGAAGAGCAACCTAAGACACATAATCGTCACATTCACCAATGTTGAAATGAAGAAAAAATG
TTAAGGGCAACCAGAGAGAAAGGTTAGGTTACCCACAAAGGAAAGCCCATCAGACTAACAGTGGATCTCTCTGCAGAAA
CCCTACAAGACAGAAGAGAGTGGGGGCCAATATTCAACTTTCTTAAAGAAAAGAATTTTCAACCCAGAATTTTCATATCC
AGCCAACTAAGCTTCAAAGTGAAAGAAGAAATAAAATCCTTTACAGACAAGCAAATGCTGAGAGATTTTGTACCACC
AGGCTTGCTTACAAGAGCTCCTAAAGGAAGCACTAAACATGGAAAGGAAAAACCAGTACCAGCCACTGCAAAAACATA
CCAAATTGTAAAGACCATCAACACTATGAAGAACTGCATCAACAGGCCAAAATAACCAGCTAGCATCATAATGACAGAA
TCAAATTCACACATAACAATATTAACCTTACATGTAAATGGACTAAATGCCCCAATTAAAAGAAACAGACTGGCAAATT
AGATAGAGTCAAGAAGCAACGGTGTGCTGTATTTCAGGAGACCGATCTCACGTGCAAAGACACACATAGACTCAAAATAA
AGGATTGGAGGAATATTTACCAAGTAAATGGAAAGCAAAAAAAGCAGGGGTTGCAATCCTAATATCTGATAAAACAGAC
TTTAAACCAACAAAGATCAAAAAAGACAAAGAAGGGCTTTACATAATGGTAAAGGGATTGATGCAACAAAAAGAGCTAA
CTATGCTAACTCTCCTAAATATATATGCACCCAATACAGGACCACCCAGATTCAAAGGCAAGTTATTAGAGACCCACA
AAGAGACTTCAACTCCCATAACAATAAGTGGGAGACTTTAACGACCCACTGTCAATATTAGACAGATCAATGGGACAG
AAAATTAATAAGGATATTTCAGGACTTGAAGTCTGAGTCTGGACCAAGCAGACCTAATAGACATCTACAGAACTCTCCACC
CCAAATTCATAGAATATACATTTCTTCTCAGCACCACATCACACTTATYCTAAAATTGACCAACAGAATTGGAAGTAAA
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ATTAAGAACTGACCCAAAACCTGCACAACCTACATGGAAATTGAACAACCTACTCCTGAATGACTACTGGGTAAATAATA
AAATTAAGGCAGAAAATAAGTTCTTTGAAACCAATGAGAACAAAGGCACAACATACCAGAATCTCTGGTACATACCCAA
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TCACAATGAAAAGAAGCTAGAGAAGCGAGAGCAAACACATTCAAAAGCTAACAGAAGACAAGAAATAACTAAGATCAGAG
CAGAACTGAAGTAAACGGGAGACACGAAAAATCCTTCAAAAAAATAATCAATGAACCCACGAGCTGGTTTTTTGAAA
GATCAACAAAATAGACCACTAGCCAGACTAATAAAGAAGAAAACAGAGAAGAATCAAATAGACACAATAAAAAATGATA
AAGGGGATATCACCCTGGTCCACAGAAATATAAACTACCATCAGAGAATACTATAAACACCTCTCTGCAATAAATT
AGAAAATCTAAAAGAAATGGATAAATTCCTGGACACATACATACACCTCCCAAATCTAAATCAGGAAGAAGTTGAATC
CCTGAAGAGACCAATAACAAGTTCTGAAATTGAGACAGCAATTAATAGCCTACCAACCAGAAAAAGTCCAGGACCAAAC
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AGGGACTCCTACCTAACTCAATTTATGAGGCCAGCATCTTCTGATACCACAACCTGGCAGAGACACACACACACAC
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CAGCACATCAAAAACCTTATCCACCATGATCAAGTAGGCTTCATCTCTGGGATGCAAGGCTAGTTCAACATATGCAAAAT
CAATAAACATAATCCATCATATAAACAGAACCAATGACAAAACCGCATGATTATCTCAACAGATGCAGAAAAAGCCTT
CGATAAAATTC AACACCCCTTCACGCTAAAACTCTCAATAAACTAGGTATTGATGGAAGGTATCTCAAAATAATAAGA
GCTATTTATGACAAAACCCACAGCCAATGTCATACTGAATGGGCAAAAGCTGGAAGCTTTCCCTTTGAAAACCCAGAACAA
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AATAAGAGAGGACACAAATGGAAAAGCATTCCATRCTCATGGATAGGACGAATCAATATCATGAAAATGGCAAAATGGC
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TAAAACTCTAGAAGAAAACCTAGGCAATACCATTTCAGGACATAGGCATGGGCAAAGATTTTCATGACTAAAACACCAA
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CAAAGAAGACATTTATGGAGCCAACAAACATATGAAAAAAGCTCTTCATCACTGGTCATTAGAGAAATGCAAAATCAA
ACCACAACGAGATACCATCTCATGCCAGTTAGAATAATGATCATTAATAAAGTCAGGAAACAACAGATGCTGGAGAGGAT
GTGGAGAAACAGGAACACTTTTACACTGTTGGTGGGAGTGTAATTAGTTAAACCATTGTGCAAGACAGTGTGGCAATT
CCTCAAGGATCTAGAACAAGAAATACCATTTGACCCAGCAATCCCATAACTGGGTATATACCCAAAGGATTATAAATCA
TTCAACTATAAAGACACATGCACACGTATGTTTATTGCAGCACTGTTTCACAATAGCAAAGACTTGGAACCAACACAAAT

Fig. 9.348

GCCCACCAAGGATAGACTGGATAAAGAAAATGTGGCACATATACACCATGGAATACTATGCAGCCATAAAAAGGATGAG
TTCATGTCCTTTGCAGGGACATGGATGAAGCTGGAAACCATCATTCTCAGCAAACACAAGAACAGAAAACCAAACACTG
CATATTCTCACTCATATGTGGGAGTTGAACAATGAGAACACATGGACACAGGGAGGGGAACATCACACACTGGGGACTG
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ACCACCATGGCATGTGTATACCTATGTAACAAACCTGCACGTTCTGCACATGTATCTCAGAACTTAAAGTATAATAAAA
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TTCATTCTCTCACCAGCAATGGCATTTGGGTTTTTTTAAAGGCCACTTTAAAAGACATTGAAAATAATACCTGTTTGTT
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TATTTATGGGATACATGTAGTATTTTGTACATGCACAGAACATGTAATGATCAAGTCAGGCTATTTGGGCTATTCATC
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TACTATAGTCACCCTACTCTGTTATTGCATATTAGAATTTTTTTCCTTCTGTGTGTTTGTACCCATTAACCAACCTCTAC
TTCATTCACCCCCCACCACCCACACACCTTCCAATCCTCTGGTGTCTATCATTCTATTCTCTACTTCCATAAGATCCA
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CCCTTCAGTTCTATCCAAGTTGCTGTAAATGCCATCATGTTAGTCTTGTTTTATGGCTGAATAGTATTCCATTGTGTATA
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ATAACATGGGCTGCAGGTATTCCTTTGATATATTAATTTCTTTCTTTGGATAAATACTAGTTAGATTACTGGACTG
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CTCCCACCAACAAGAATTCCTTTTTTCTGCATCCTCACCAGCATCTATATTTTCGTCTTCATCTTCTCCTTCTCCTTCT
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TCAAGCAATTCTCCTGCCTCCCCTCCCAGTGGCTGGGATTACAGGCGCCACCACCATGCCAGCTAATTTTTGTATT
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CCCACAGTGCTGGGATTACAGGTGTAAGCCACTGCACCTGGCCCTTTACCCACTTTTTAGTAGGATGATTTGTGGTCTT
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ATTTTTCTTTTTGGTGCTGTGATCTAGAGATCTTAGCTATAAAATCTTTGGTCAGACTGATGTCCTGAATTGTTTTCC
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CCGATGGTATATATAGGCACTAGCTATGGTAGGCAGGGGCATGGTGATTTCCAGGCCCTCAGTGGAATGCTTGGATGGG
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GGAGGTCAATGGGGCTCAAGGAATCTGGAGTTGCAAGGTCTGTGGGGTCCCAGGGTAGGATGCAGTCTGCTGGGCTTTC
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ATGTTTAAAGGCTAGTTGAATAGGGGTGAACACAGAAAATCCTAGCAATTCATTCTTTGCACTCCTGGCATTCTCAAATG
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TGAAATAATTCTGACAATAATTAGTATTTTTTTATAGGAATTGGAACATGGTTTATTGCAGTATACCCAGTGTTTGTCA
GCTCTAGAACTTACGAAGAAGAGGAAGAGGAGAACAAGTTTGACTCCTGTGAACATTTTCTTTTTTCTATAATTTACG
CATATGCTAACAAGCAGCTAGTTTCAAACACAAACTTACCTTGACAGGAACTAGGTCTCACTGTTGAGAATGTAGTCT
CCTCTCTCCCCACATAGCCCTAGGGTTATTTCAGGATGAAGGAGAAGAAAAACATACCCAAGAGCATTTTATAATTCCA
TCCTTCCTTCTTGTATGCTAAGGTTACTAGTACCATGACCTATGTGAACCTCGTTTTCTTGAATAAGAAGAATAAAAGC

Fig. 9.349

GTTCCGTC CATCAAGGAAGACCTCAAGAGAAATTCCAGGTT CAGGTCTCATGGTACAAGAGCCAAGTGTTTCTTCCTGA
ATCATTTCCAGGCCTGCTTTTGATCTGAGCAGTGGCTTTCAAAAAATATGCTGATGAGATTGCTCTCTCACTTAAGAAC
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GTCCACTGACATGCCCTTGGA CTCCAGGAGAAATCTGGAAATGTCTGGACCTGCAATGAGACACACCCAGCTTCAAAT
GTGTC ACTCTGGGCATATTCTTTAATCTGAGCTCCAATTTTCTTTCTTGTAAGAGTCATTACACATGGCTGTTTTATTT
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GGCTCATTTCCATCCATTTTCATGGATGTGTTACCTTCTGGTATTTATTAGAATTTGCTTTTAGACTTTCTGTTGCTCACC
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CTCCTGTTTTTTGAAACCCACCAGTGTCCCCACTTCCATAAGTCCCTGGATCCTCAATTTTCCAAGTCTTCCTGGAATTC
TATGGAGTAAATTAGCTTACTGATGCATTTCCCTTACTTGCTTAGGTTTCTGCTCACTCTGTTCCCAAGTCAATTATG
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CATTTTAAAAAATTCTTTTGCTGTCGTTTTTGTGGAGAAATAGAAAAATATGTACATTCAATCCCCATGTTTAAGTGGA
AGTCCCTCATAAAATTATATTAGAAATTTATTAGAAATATATACACACATATGAATATAGATAGATATACACTCTCTTT
TTTAGCATACATAGTGCCTGTTATTTAGCAGGTACTAAAAAATAATATGTATATGTATATACATGTAGCAGAAGGCTAA
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GACTTAGAAAAAGACGGAACCATGGCTGGGTGCGGAGGCTCATACTGTAATCCCAGCACTTTGGGAGGCCGAGGCCAG
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ATTGATTTTTTACAAGCTAGCAAATATTTTCAATAGTTGAAGCTTCGGAGTTTCTCATTATCTAAGCTATGTAAATGCA
TGCAAAGTTTCTTCTAGAAAACAAATTA CTGAAAGACATTTTCCTGATTTGTATTTGGCTGTGTTATTCCCCAGAAGGT
GAAATTATTAAACATGCCATTTCGAAAGCCAGTAACTCCTTAGTACAGGTTGAACATCCCTAATCTGAAAATCTGAAATC
TGAAATGCTCCAAAATCTGAAACTTTTTCAATGCCAACATGATGCCACAAGTAGAAAATTTTACACCTGATCTCATGTG
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GGAAATAAAAAATAAACTCTAAGTCCCTCCGACTGACCCAGCGGATTCTCTCTTGGCCAAGGGAACCCAGCAAAACCT
TGGAAGCTGAATTCATGGCTATGATGGGATGGGAGATTTGGCATATGCCTCATTATATCCCCACCCTCGCTAACAGTCG
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GTCCAGATGAGATTCAATTATGTTTACCTTATTTTATGTAAGATGTAGATTTACCAGGCACTAACTAAAGTTTTACAAGT
ATGTAATCATTTGTCTCACTGCTGCCCCACCACCTCCCCTGCCTTTTAAAGGAAAATATATAAATACTAAACCTCCTAAG
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CTGGCAGCTGAAAATCTGAGTCACTATCATGTCAA ACTATAAAAAATTTTATACA ACTTAAAAA AACTGC ACTAA
TCATTAATTTATATTATTGTTAAATTTATCATACAGTAAATTTGATTTTTTTTCTTTTGATGTACAGTTCTGAAACCA
CCTTTGCAAAAATTGCAACACTGAGAAA ACTTTGACAGTGAAAGAAATTTGACCTAACCAACTCCACATTGCCTTTAAC
CTCCAAACTGCCCTTCATTCTGGGCATGGCCTAAGCTAACTTTGGGAGAAATTTAGGTTATAGTTTAAATGATAATAG
CTCTTTCCAAA ACTAACTGCCTTTGTAAA ACTAATGAAAGGCCACCAGTTTACGAAGATAGGAGGGCCTGAATTCTGC
TAAGATATAGGCATAGTTAAGTGATTACCAGCCATTATTCAGAGGTCACAAGATTTTCAACTTCTCAATTACTCCTG
TAAATAACGTTACTATTGTAGAACCTAAAATTACTATTGTAGAACCTAAAGTTGACCTTTTGAGATGTCTTGT CAGGCT

Fig. 9.350

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TTTGCATTTCTGATGACMCCAGTGTCTCTGAACCAGTGACTCCTCTGTGGACCCTTACTGGAAGCTGACTCAGGGCACAC
GAGGACCATTTTCCACACCCATATGATTGCATCCCCAACCAATCAGCAGCACCCATTCTTTGCCCACCAATTACTCT
TGAAAACTCTAGCCTCCAAATTTTCAGGGAGGCTGATTTGGGTAATAATAAACTCTGGTCTCCTGTTTAGCTGGCTC
TATTTGTATTAACTCTTTCTCTACTGCAATTGCCCTATCTTGATAAATCAGCTTTATCTGAGCAGCAGGCAAGAAGAA
CCCATTAGACAGTTACAGTTCTGTAAATTTTAAACACAAACATTGTTTCATGTAACCACCAGTAGGATCACCATCTGGATC
AGTTTCACCACCTAAAAAGCTTCTCATAATACCCTACTCCATGACACTAGGAACAGCTACTAGTTCTCCATCCCAATT
GTTTCATCTTTTGAGAAGTTTATATAAATGGAATTGTATAGTATAATCATCAATTTTAAATAAAATTTTCAGATGTATT
CACATTCCATAAGTGTACTCTATGATGCTGATATTTATCTTTAGCCTCAGCTTATGCCAAACTATGCCTTTCCATATAT
GGCCAAAATGTGTATTCAATATACTCTAACTGGGAGGATCAATTGAAAGTGTTGTTTTACTAACTCTATAACACAAA
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TGGGTGGTTTTATAACAACGAACATTTATTTCTCACAGTTCTGGAGGCTGAGAAGTCCGAGATCATGCTGCCAGCAGAT
TCACTGTCTGGTGTGGTCCCATCCCTACACAGTGAAAGCAGCAAGGTCACTTTCTGTGCCTGATTTATAAGGACACTAA
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TTCATTCTGATGCCCAGGTATGACAGACCTCAAATCTGTACCGTCCCATGAACCAGGAATTATACAAAGGGTAGAGGG
GCATAGCTCTGGGTGAAGAGCTGCACTCTGTTCACTCTGGGCTCTGTGTGGTGGTGTCCCAGGTAGTGTCACTCCAGCT
GTTAAGTCCAGTGAGGCTCAGAAAGGTGATGTCAATTTGTTCAAATCTTCCAGCTAGTTTGTGGTTCAGATCATAACCAG
ACCCTATGAAGTCTAATCATAACCAGTTCTGTAGGCTTTCTCTTCAATAGGTCCATTCTTATCTCTAACTGAGACCAC
CTGCATCCTCATAGCTTAACTATCAGTATGGAAAACAAACCATCTGTGTCACTATGACTTTACTATTTGTTCAAGAAAA
TTCTTGTCCAGGAAGATAAGACGGTGTACAATAAATAACATCAATGTTTACTTCAGAAAAATTTCTGAAAACCAATTTAT
TCAGATGATAGATTGCTCATTTGAAAAAATAAACCCCTTTTCAGGACAACAGATTCCTCTCCAAGTCTAATAACTTGT
TATCAAAGATCTATTTTTCAGGACTTCAAGACCTCTTATCACAATGTCCACCAATTCTAAACTATAATATCATGAAC
TTGCCGAATTTCCACCAGATTTCTTCTTGAAAGGCCTGACTTTAAACCACTTGAGCTCAGACCCTTAATGTTTATAAA
TATCTACCTGTGACCTCTCCCTTTTGAGAATTATAAGGACTTTTTCAAGGTGTTGCACTCTCTTACTGCAGGTTAATAA
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GGATTCACCTCAATAATTTCTGCTCCTGCAGTTTAAAGAACCTTAGCCCTAAACAGTGTGTTCTTCTGAAGTTTGCCATTT
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CTTTGTGGAAATGTTTATTACTAGTAACATTACTCTTTTGTCTTTTATCCTATTTGTTGTGAGTCTATGAGAAGAAG
CTCACAGGGAAGAAGACAGCCGTAGACCTGGCAATTTTGTCTCAAACTGGCTCAAAGATAACAATAATATCAGCGGGTC
TCATCAAACCTGGCATTCTTATAGAATAAACTTTGCTCTGGGTCACTTACTAATATCTTACAGAAAATTTATAGCAGCAG
TTGTATATTGAGGGTGTGAATAAAAACCACCAGAGAAGCTTTTTGAAAAAATACTATGAATAGATTTCTGCTTCTGGCC
AAAATGAAGAAACAGGGACCAGATTTACTCTCCAGTAAACCAGTGGAAAAAAAACAGACACAATATTGAAAAATAAAA
AGATTTTTCAGACATCAGGCAATGAAGAATAGTGATCCAAGAGAAACAAGAAACAAGATGAATCTTATGTTTCGCCACA
GCTTACTGTCTGGAGTGAGTATCAAGACTGTGGTACAGAAAGGGAAAACCCAGATGGAACCCTGCCATCTCCCTAAGTG
GAGTCTGGGGAGGGCAAAGTGAGTAGAGTTTGCAAGGAAAGATATGGGAAAGGAGACAGCTGTGCAGAGAGAACACTGG
GGATCTGTACAGGGTCTTCAGCTGAGCATGAGTCAGCATATGCATGTGAGGAACTACCCATGACTGGGGAAAGAATCA
GCTGGAATGATTACGGGGTATAGAATCCAGGGCTCAGAATCATTCCTGTTTCTATAGGAAATGGGCATTGAGTAGCATA
CTTGGAAGAATTTTGTCTTCAGTCATGCAGTAACATAGAATCTAGACTAAATACTACTCTGATACCACATAATGAACT
CAAAATAAGACCCAAAAGAATCAAACCTGTTTATAAGTAACCTGAGTTTCAAAATAAAGTTCAAGAATATTTATAGAAATA
CCATCTAACAAAGCACAAAATTAAGATAAAGGGTCAAACCTATTTGCAAGTAATTTTAAAGTGTGTTCTAGAACAAGTTT
AAAAGTATTTATAGAAATACAAAAGGATCCAAAACCTGTCCACAAAATATGACCCATAATGAAGAGAAAAATCAGTCA
TTTGAACTAACCAGAAATGACAAAGATGATAGAATCGGCAAAGACATTAGAAGAACTGTAATTGTACTTCGCATGTT
CAAGAAGCCAGAGGAAAGACTGAACATGGTAAGCAGAAACATGGAAGATATAAAAAGACTAAAAATCAAACCTTTAGAGA
TGAAAACATTATGTGAGATGAAAAACACACTGAGTAGAATTAAAGGCAAATTGGAAATTTTCAAGAACTAGTGACTTTA
AGAATTGAGAGATAAAAAGTACACAAAATGAGAAACAGAAGTGTGTCACTGAGTTATGGGACAACCTTCAAACCTAATGCG
TAAATTGCAGTCCCTGAAGGAAATGAGGGATATGTTGGAAAAAATATTTGAAAAAATAATGGCCAAAAATCTCCCAA
GTTTATGAAAACACAGATTCAAGAAGGTCAACAAATTGCTTAAAAATAGAGAAAGTCTAATATTAGGATACAAGGTCT
TGACCAGGTGTGGTGGTTCATGCCTGTAATCCCAGCACTTTGGAAGGCCGAGTGGGAGGTGAATCACTTGAGGTGAGGA
GTTCAAGACCAGCCTGGCCAACATGGTGAAAGCCTGTCTCTACTAAAAACACAAAAATTAGCCCGGCGTGGTGGTGCAC
ACTATAATCCCAGCTACTCGGGAGGCTGAGACAGGAGAATCACTTGAACACAGGAGGTGCAGGTTCAGTGAGCCGAGA
TTTGTGCATTGCACTCCAGCCCTGGGTGACAAAGTGAGACTCCGTCTTAAAAAAGACAAAAACAAAAACAAAA
AAATGGAAATAAGGTCTCAAATTAATAACTTCAGCTTACACCTTAAAAAAAATTAGAAATATCATT

Fig. 9.351

Single marker association within the PDE4D gene



FIG. 10.1

FIG. 10.2

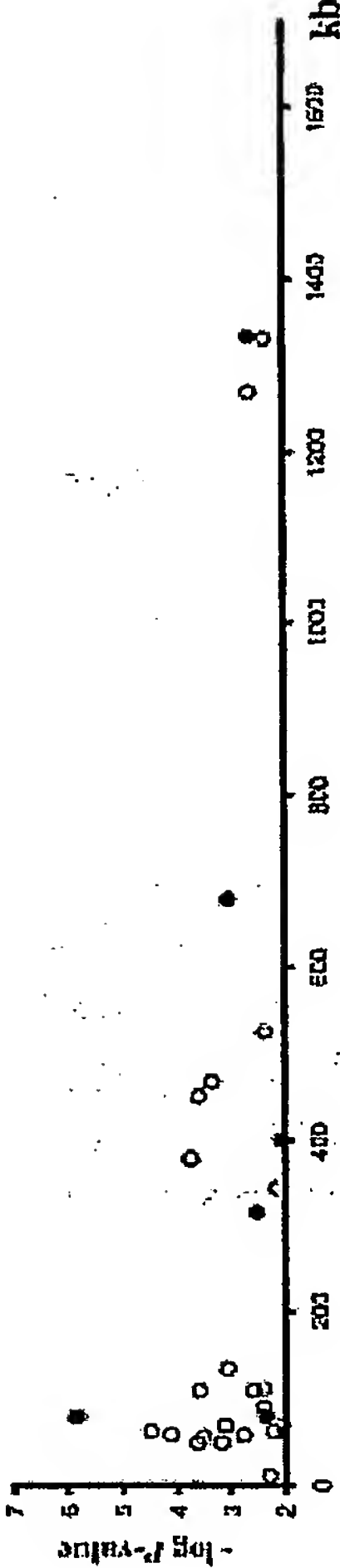


FIG. 10.3

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FIG. 11.1

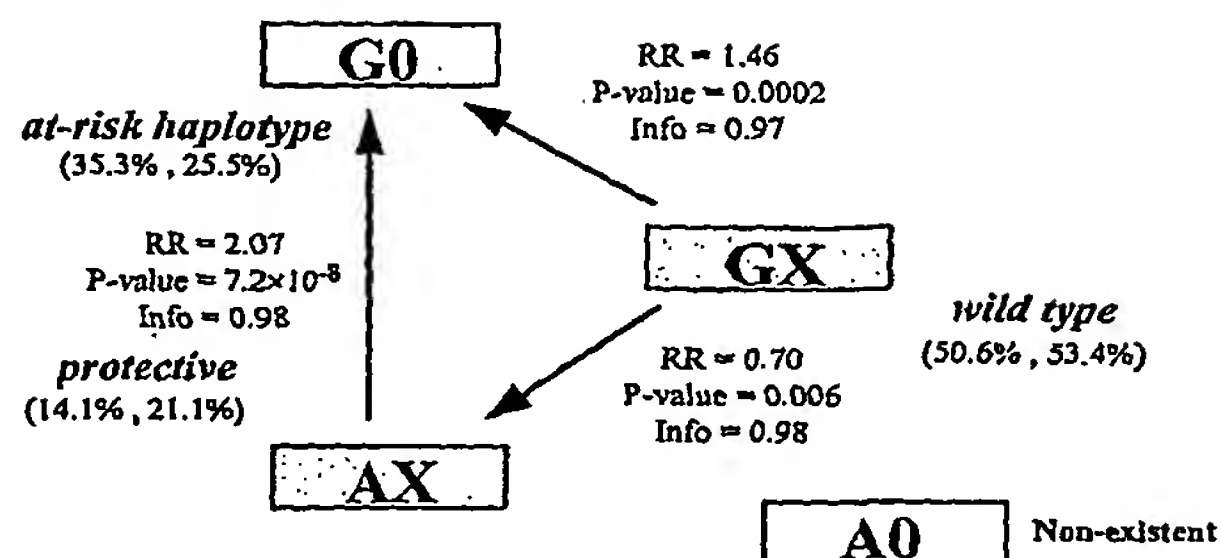


FIG. 11.2

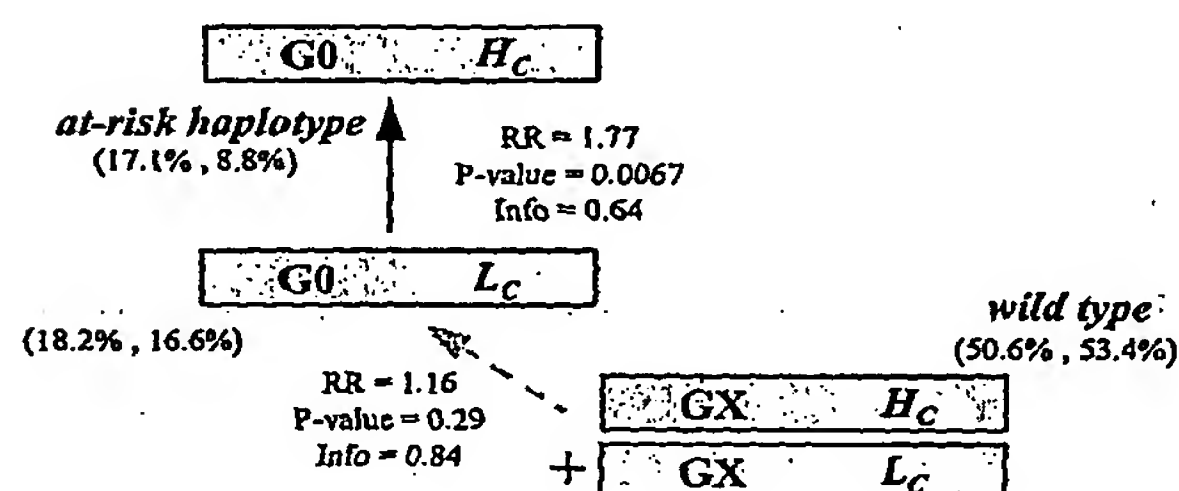
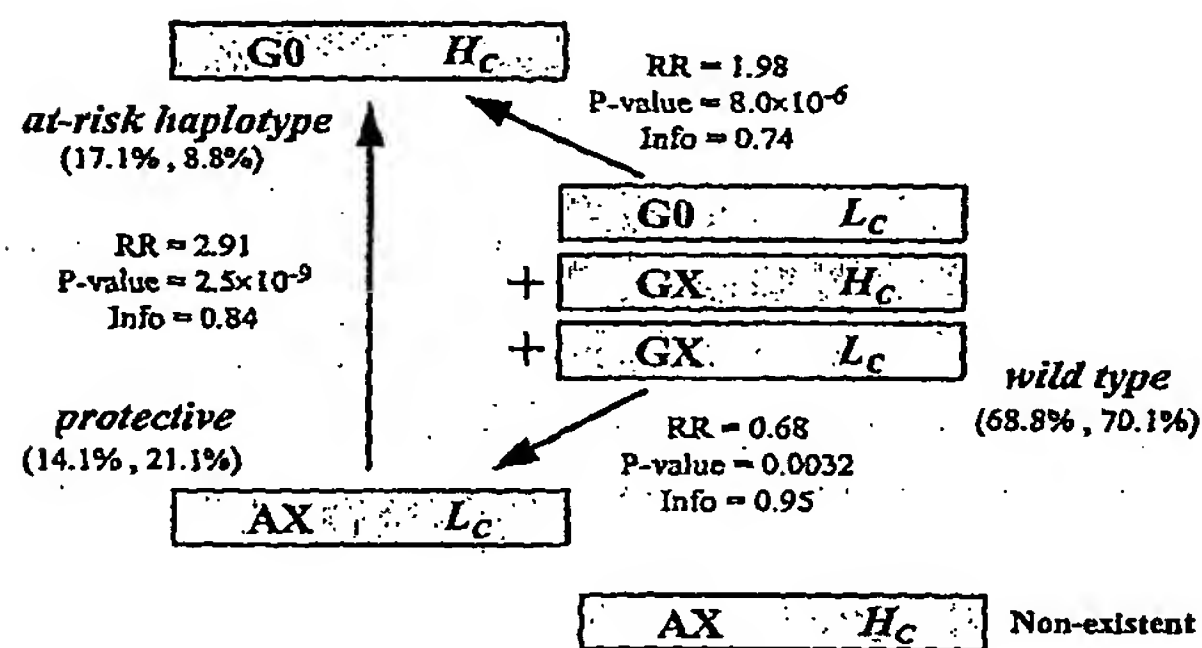


FIG. 11.3



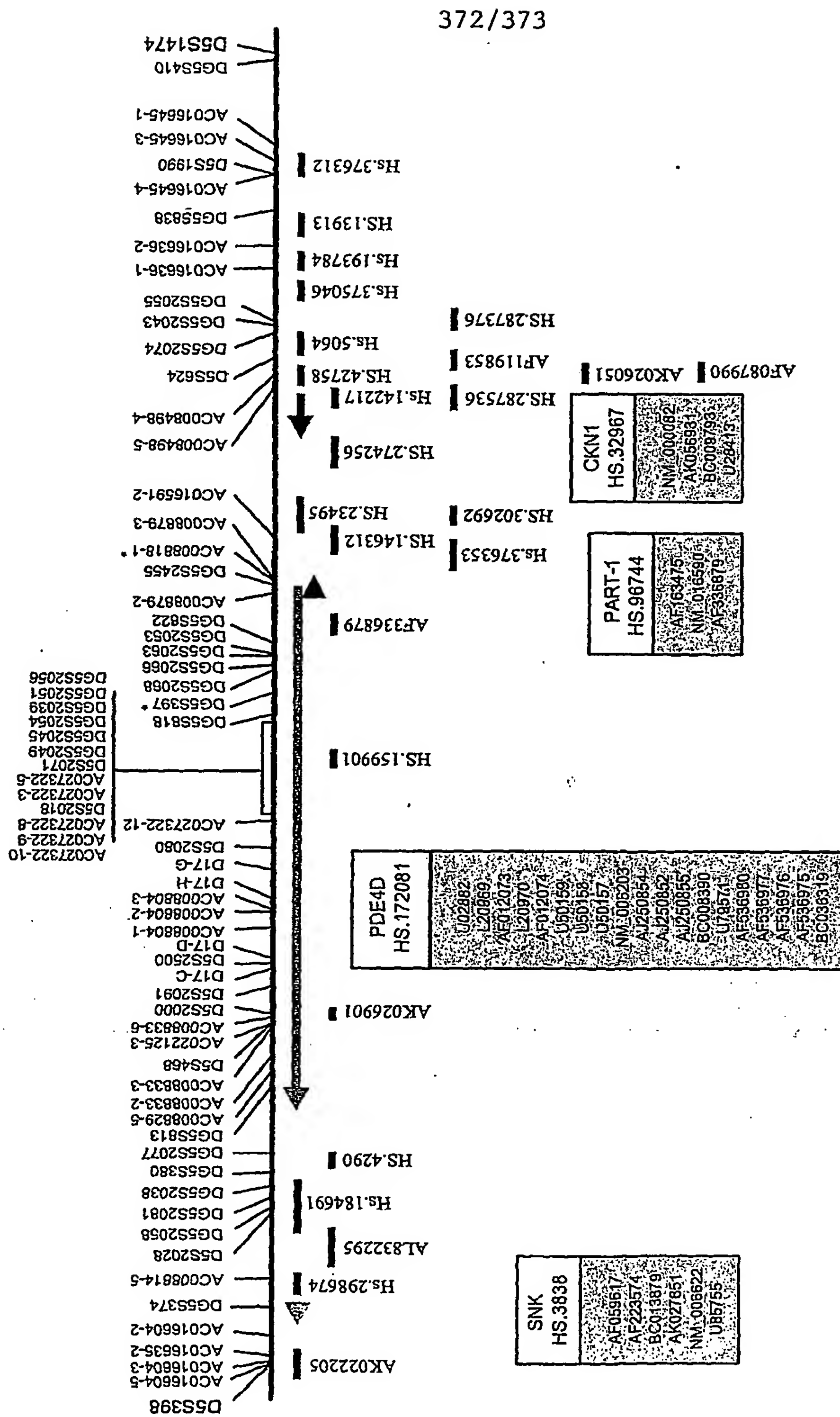


FIG. 12

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FIG. 13.1

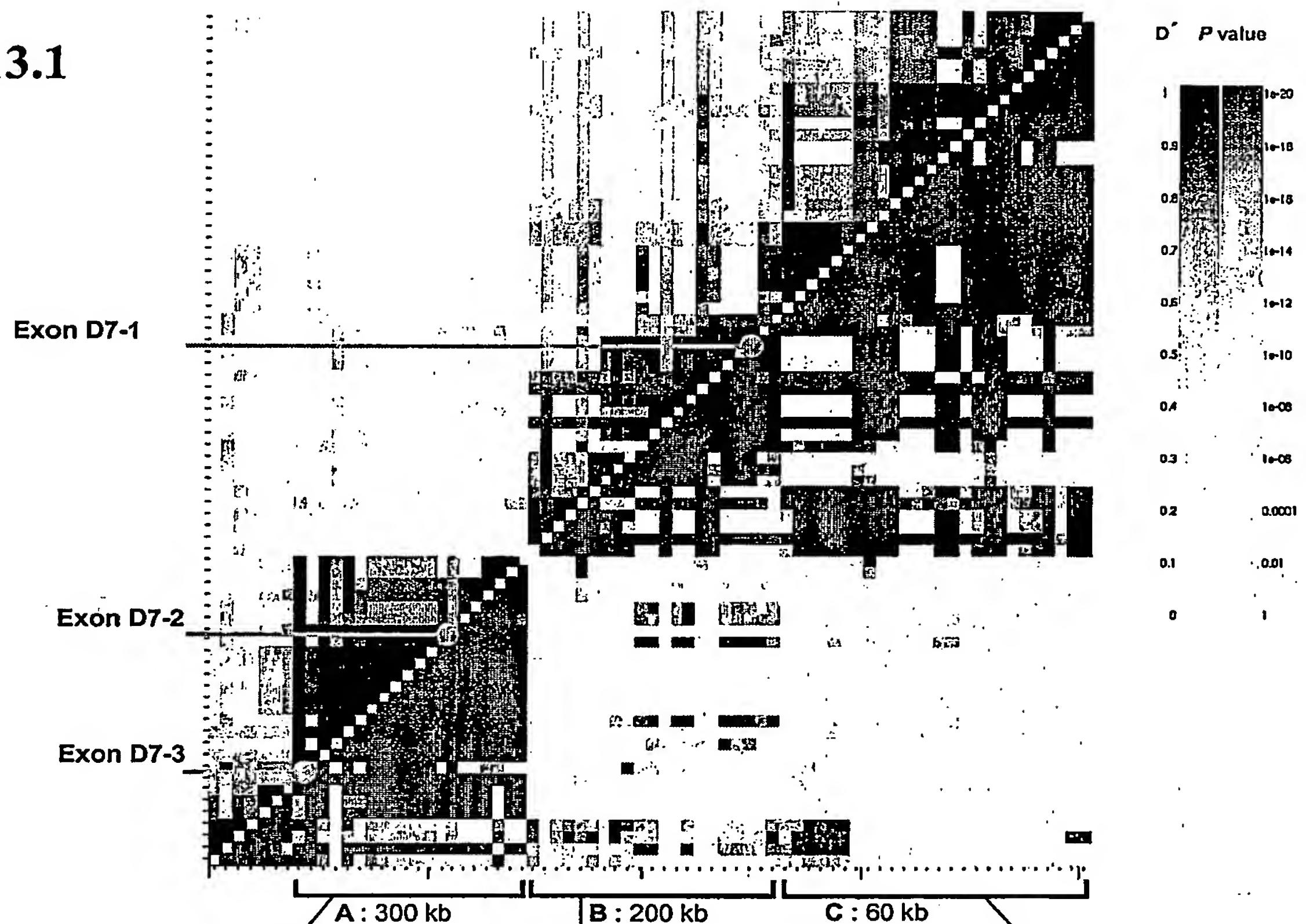


FIG. 13.2

TGTATGAGAGAGTAGCGGTC
 CGCACCGAGAGACGGCGGTC
 CGCACCGAGAGACGATGGAT
 CGCACCGAGAGACGATAGAT
 CGCGCGAGAGAGTAGCGAAT
 CACACCAGAGAGTAGCGAAT
 CACACCAGAGAGCAATGGTC
 CACGCGAGAGAGTAGCGAAT

AATGTAAGAACAGTACCTGAAT
 AATGTAAGACTAAAATTCAGGA
 GACATAAGAACAGTACCTGAAT
 GACATGGAGATAAAAATTCGGAT
 GACATGGAGCTAAAATTCAGGA
 GGCATGAGAACCCTGTCTGAAT
 GGCAGGAGAACCCTGTCTGAAT

TAACCACGAACTTATTGAATTTGAA
 GAACCACGAATCCGCCGAGCATCAA
 GAACCACGAATCCGCCGAGTTTGAA
 GAACCACGATTCTACCAGGCACCTG
 GGCTTCCGAACTTATTGAATTTGAA
 GGCTTCCGAATCCGCTGAGCATCTG
 GGCTTCGAGTTCTACCAGGCACCTG

FIG. 13.3

84.0 %

72.9 %

70.6 %